

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: April 10, 2002, 21:32:13 ; Search time 107.54 Seconds
(without alignments)
2672.499 Million cell updates/sec

Title: US-09-380-276A-1
Perfect score: 1269
Sequence: 1 atgctttaaagtctact.....ggcagcagctgggttccctg 1369

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA.*
1: /cgn2.6/ptodata/2/ina/5A_COMB.seq.*
2: /cgn2.6/ptodata/2/ina/5B_COMB.seq.*
3: /cgn2.6/ptodata/2/ina/6A_COMB.seq.*
4: /cgn2.6/ptodata/2/ina/6B_COMB.seq.*
5: /cgn2.6/ptodata/2/ina/PCTUS_COMB.seq.*
6: /cgn2.6/ptodata/2/ina/backfiles1.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	435.4	34.3	893	4	US-09-286-529-8
2	314	24.7	623	4	US-09-286-529-9
3	36.4	2.9	1601	1	US-08-722-001-7
4	36.4	2.9	1987	1	US-08-722-001-26
5	36.4	2.9	1997	1	US-08-722-001-27
6	36.4	2.9	2004	1	US-08-722-001-11
7	36.2	2.9	2485	1	US-08-424-424B-1
8	36.2	2.9	2486	5	PCT-US94-05363A-1
9	36	2.8	4136	4	US-09-103-875-2
10	35.6	2.8	1150	4	US-09-372-934-3
11	34.8	2.7	1639	1	US-08-334-698-5
12	34.8	2.7	1639	1	US-08-228-932-5
13	34.8	2.7	1639	1	US-08-468-939-5
14	34.8	2.7	1639	1	US-08-406-855A-5
15	34.8	2.7	1639	2	US-08-722-190-5
16	34.8	2.7	1639	3	US-08-244-354-5
17	34.8	2.7	1639	3	US-09-206-899-5
18	34.8	2.7	1639	5	PCT-US95-04203-5
19	34.6	2.7	5962	6	US95-04203-5
20	34.6	2.7	7218	1	US-08-232-463-14
21	33.6	2.6	800	2	US-08-416-603-11
22	33.4	2.6	4360	1	US-08-470-508B-1
23	33	2.6	9472	1	US-08-325-547-9
24	32.8	2.6	2230	1	US-08-200-512-1
25	32.6	2.6	1593	2	US-08-524-828-2
26	32.6	2.6	1593	2	US-08-975-114A-2
27	32.6	2.6	1593	3	US-08-849-281A-2

28	32.6	2.6	2247	2	US-08-524-828-1	Sequence 1, Appl
29	32.6	2.6	2247	2	US-08-975-114A-1	Sequence 1, Appl
30	32.6	2.6	3891	1	US-08-480-604A-27	Sequence 27, Appl
31	32.6	2.6	3891	2	US-08-405-496A-27	Sequence 27, Appl
32	32.6	2.6	3891	4	US-08-915-136-27	Sequence 27, Appl
33	32.4	2.6	1167	1	US-07-960-985-1	Sequence 1, Appl
34	32.4	2.6	1167	2	US-08-496-671-1	Sequence 1, Appl
35	32.4	2.6	1280	4	US-09-096-776B-4	Sequence 9, Appl
36	32.4	2.6	1491	4	US-09-082-092-9	Sequence 3, Appl
37	32.4	2.6	1524	4	US-08-840-767-3	Sequence 3, Appl
38	32.4	2.6	1690	2	US-08-461-812-3	Sequence 3, Appl
39	32.4	2.6	3083	2	US-08-480-994-36	Sequence 36, Appl
40	32.4	2.6	3083	2	US-08-616-844-36	Sequence 36, Appl
41	32.4	2.6	3083	2	US-08-599-654-36	Sequence 36, Appl
42	32.4	2.6	3083	2	US-08-485-573-36	Sequence 36, Appl
43	32.4	2.6	3083	3	US-08-944-868A-36	Sequence 36, Appl
44	32.4	2.6	3083	3	US-08-944-423A-36	Sequence 36, Appl
45	32.4	2.6	3083	3	US-08-925-743-36	Sequence 36, Appl

ALIGNMENTS

RESULT 1
US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catharine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match 34.3%; Score 435.4; DB 4; Length 893;
Best Local Similarity 82.1%; Pred. No. 1.1e-125;
Matches 513; Conservative 0; Mismatches 111; Indels 1; Gaps 1;

QY	1	atgctttaaagtctactagaacaagaaacatttttcaactcttttagttacta	60
DB	55	atggcactcaagtcctactctacacaggacggtgctctctgcca	114
QY	61	ggctattgtcattgtaaaagtgttgtaaacaggagactgtagacagcaagaattcag	120
DB	115	ctccacctggcatgtaaaagtgtgtaaacccgagattgcaggcagcaggaattcaag	174
QY	121	gacggtctggaacatgtgttcctctgcaacacagtggtggccaggatggagttgtctaag	180
DB	175	gacggtctggaacatgtgttcctctgcaacacagtggtggccaggatggagttgtccaag	234
QY	181	gaatgtggtcttggtcattgggagagatgcacagtggtgacgtgccggctgcacaggttc	240
DB	235	gaatgtggtcttggtcattgggagagatgcacagtggtgacgtgccggctgcacaggttc	294
QY	241	aaggagactggggcttcacagaaatgcagccctgtctggaactgcgcagtggtgaaaccgc	300
DB	295	aaggagactggggcttcacagaaatgcagccctgtctggaactgcgcagtggtgaaaccgc	354
QY	301	tttcagaagcaaatgtttcagccacacagtgatgcacatctcggggagactgcttgcacagga	360
DB	355	tttcagaagcaaatgtttcagccacacagtgatgcacatctcggggagactgcttgcacagga	414
QY	361	ttttatagaagacgaaactgtcggtttcaagacatgagtgatgtgcttctgtgagac	420
DB	415	ttttacgcgaagaccacacagtggtgttttcaagacatgagtgatgtgcttctgtgagac	474


```
; GENERAL INFORMATION:
; APPLICANT: Thompson, Wayne J.
; APPLICANT: Huff, Joel R.
; APPLICANT: Nerenberg, Jennie B.
; APPLICANT: Lee, Hee-Yoon
; APPLICANT: Bell, Ian M.
; TITLE OF INVENTION: ALPHAIC ADRENERGIC RECEPTOR ANTAGONISTS
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Merck & Co., Inc.
; STREET: 126 Lincoln Avenue
; CITY: Rahway
; STATE: New Jersey
; COUNTRY: United States of America
; ZIP: 07065
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/722,001
; FILING DATE:
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/229,276
; FILING DATE: 14-APR-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Appollina, Mary A.
; REGISTRATION NUMBER: 34,087
; REFERENCE/DOCKET NUMBER: 19169Y
; TELEPHONE: (908)594-3462
; TELEFAX: (908)594-4720
; TELEX: 138825
; INFORMATION FOR SEQ ID NO: 26:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1987 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: both
; TOPOLOGY: both
; MOLECULE TYPE: cDNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; US-08-722-001-26

Query Match          2.9%; Score 36.4; DB 1; Length 1987;
Best Local Similarity 52.7%; Pred. NO. 0.27;
Matches 79; Conservative 0; Mismatches 71; Indels 0; Gaps 0;

QY 474 cgcgtccacggcctccagccacgggacacggcgctggctgcgttatctgcagcgctct 533
DB 1112 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTGCTCTTCGGCTCT 1171
QY 534 ggccaccgctcgtggccctgctcatctctgttcattctattgtaagagacagtttat 593
DB 1172 GGGCTCTTCTACCTGCTCTGGCCATCATCTGGTGCATGTACTGCGGCTCTACGTGCT 1231
QY 594 ggagagaagaaacccagctggtctctcggtc 623
DB 1232 GGCCAAGAGGGAGAGCGGGGCTCAAGTC 1261

RESULT 5
US-08-722-001-27
; Sequence 27, Application US/08/722001
; Patent No. 5760054
; GENERAL INFORMATION:
; APPLICANT: Thompson, Wayne J.
; APPLICANT: Huff, Joel R.
; APPLICANT: Nerenberg, Jennie B.
; APPLICANT: Lee, Hee-Yoon
```

```
; APPLICANT: Bell, Ian M.
; TITLE OF INVENTION: ALPHAIC ADRENERGIC RECEPTOR ANTAGONISTS
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Merck & Co., Inc.
; STREET: 126 Lincoln Avenue
; CITY: Rahway
; STATE: New Jersey
; COUNTRY: United States of America
; ZIP: 07065
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/722,001
; FILING DATE:
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/229,276
; FILING DATE: 14-APR-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Appollina, Mary A.
; REGISTRATION NUMBER: 34,087
; REFERENCE/DOCKET NUMBER: 19169Y
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (908)594-3462
; TELEFAX: (908)594-4720
; TELEX: 138825
; INFORMATION FOR SEQ ID NO: 27:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1997 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: both
; TOPOLOGY: both
; MOLECULE TYPE: cDNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; US-08-722-001-27

Query Match          2.9%; Score 36.4; DB 1; Length 1997;
Best Local Similarity 52.7%; Pred. NO. 0.27;
Matches 79; Conservative 0; Mismatches 71; Indels 0; Gaps 0;

QY 474 cgcgtccacggcctccagccacgggacacggcgctggctgcgttatctgcagcgctct 533
DB 1106 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTGCTCTTCGGCTCT 1165
QY 534 ggccaccgctcgtggccctgctcatctctgttcattctattgtaagagacagtttat 593
DB 1166 GGGCTCTTCTACCTGCTCTGGCCATCATCTGGTGCATGTACTGCGGCTCTACGTGCT 1225
QY 594 ggagagaagaaacccagctggtctctcggtc 623
DB 1226 GGCCAAGAGGGAGAGCGGGGCTCAAGTC 1255

RESULT 6
US-08-722-001-11
; Sequence 11, Application US/08/722001
; Patent No. 5760054
; GENERAL INFORMATION:
; APPLICANT: Thompson, Wayne J.
; APPLICANT: Huff, Joel R.
; APPLICANT: Nerenberg, Jennie B.
; APPLICANT: Lee, Hee-Yoon
; APPLICANT: Bell, Ian M.
; TITLE OF INVENTION: ALPHAIC ADRENERGIC RECEPTOR ANTAGONISTS
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Merck & Co., Inc.
```

STREET: 126 Lincoln Avenue
CITY: Rahway
STATE: New Jersey
COUNTRY: United States of America
ZIP: 07065
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,001
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/229,276
FILING DATE: 14-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Appollinis, Mary A.
REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19169Y
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)594-3462
TELEFAX: (908)594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 2004 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-722-001-11

	Query Match	2.9%	Score 36.4	DB 1	Length 2004	
	Best Local Similarity	52.7%	Pred. No. 0.27			
	Matches	79	Conservative	0	Mismatches	71
					Indels	0
					Gaps	0
QY	474	cgcgctccacggcctccagcccccagggacacggcgctcggtccggttatctcagcagctct	533			
Db	1107	CGAGGACGAGACCATCTGCCAGATCAACGAGGACCGGGCTACGTGCTTCTTCGGCTCT	1166			
QY	534	ggccacacgctcgtcggtggccctgctcctctctgctcatctatgtgaagacagagttat	593			
Db	1167	GGGCTCCTTCTACCTGGCTCTGGCCATCATCTGGTCTACTGTACGCCGCTCTACGTGGT	1256			
QY	594	ggagaagaaccccgctggtctctctgcggtc	623			
Db	1227	GGCCACAGGGGAGAGCCGGGGCCCTCAAGTC	1256			

```

RESULT      7
US-08-424-424B-1
: Sequence 1, Application US/08424424B
: Patent No. 5759854
: GENERAL INFORMATION:
: APPLICANT: LI, ET AL.
: TITLE OF INVENTION: Neurotransmitter Transporter
: NUMBER OF SEQUENCES: 6
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
: ADDRESSEE: CECCHI, STEWART & OLSTEIN
: STREET: 6 BECKER FARM ROAD
: CITY: ROSELAND
: STATE: NEW JERSEY
: COUNTRY: USA
: ZIP: 07068
: COMPUTER READABLE FORM:
: MEDIUM TYPE: 3.5 INCH DISKETTE
: COMPUTER: IBM PS/2

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; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WORD PERFECT 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/424,424B
; FILING DATE: APRIL 21, 1995
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/US94/05363
; FILING DATE: MAY 25, 1996
; ATTORNEY/AGENT INFORMATION:
; NAME: MULLINS, J.G.
; REGISTRATION NUMBER: 33,073
; REFERENCE/DOCKET NUMBER: 325800-308
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 201-994-1700
; TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2485 BASE PAIRS
; TYPE: NUCLEIC ACID
; STRANDEDNESS: SINGLE
; TOPOLOGY: LINEAR
; MOLECULE TYPE: cDNA
; US-08-424-424B-1

Query Match 2.9%; Score 36.2; DB 1; Length 2485;
Best Local Similarity 56.2%; Pred. No. 0.36;
Matches 68; Conservative 0; Mismatches 53; Indels 0; Gaps 0;

QY 453 caagggtcaacctgtggaagatcgctccacggcctccagccacgggacacggcgctggc 512
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Db 2023 CAGCATCATCCAGTGGGGGTACGCCCGCGGCTACAGCGCTGGATCAAGGAGGAGGC 2082

QY 513 tgcgcttatctgcagcgctgtgcacacgtctctgtgctgcctgtcatctctgtgtcat 572
   ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db 2083 TGCCGAGCGGTACTCTGTATTTCCCAACTGGCCCATGGCACTCTCTGTATCACCCCTCATCGT 2142

QY 573 c 573
Db 2143 C 2143

RESULT 8
PCR-US94-05363A-1
; Sequence 1, Application PC/TUS9405363A
; GENERAL INFORMATION:
; APPLICANT: LI, ET AL.
; TITLE OF INVENTION: Neurotransmitter Transporter
; NUMBER OF SEQUENCES: 2
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
; ADDRESSEE: CECCHI, STEWART & OLSTEIN
; STREET: 6 BECKER FARM ROAD
; CITY: ROSELAND
; STATE: NEW JERSEY
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 INCH DISKETTE
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WORD PERFECT 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US94/05363A
; FILING DATE: SUBMITTED HERewith
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: FERRARO, GREGORY D.
; REGISTRATION NUMBER: 36,134

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COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/98/228,932
FILING DATE: 13-APR-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41878-B/

APPLICANT: Jonathan A. Bard et al.
TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic Receptors and Uses Thereof
INVENTION: 6
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/468,939
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41337-1B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEFAX: (212) 391-0526
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1639 base pairs
TYPE: nucleic acid

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; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHETICAL: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523
; OTHER INFORMATION:
; US-08-468-939-5

Query Match          2.7%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 474 cgcctccagcgcctccagccacggcgcgtggtgcgttatctgcagcgtct 533
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTTCTTCAGCGCT 694

QY 534 ggcacccgtctgtggtgcctctcatctctgtgtcatttgaagagacagtttat 593
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 695 GGGCTCCTTCTACCTGCCTCTGCCATCATCTGCTGATGCTACTGCCGCTACGTTG 754

QY 594 ggaagaagaacccagctggtctcgcgtc 623
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DB 755 GCCAAGAGGGAGAGCGGGGCTCAAGTC 784

RESULT 14
US-08-406-855A-5
; Sequence 5, Application US/08406855A
; Patent No. 5861309
; GENERAL INFORMATION:
; APPLICANT: Jonathan A. Bard et al.
; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
; TITLE OF INVENTION: Receptors and Uses Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Cooper & Dunham LLP
; STREET: 1185 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/406.855A
; FILING DATE: 21-AUG-1995
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41337-A-PCT-US/JPW/KDB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0526
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHETICAL: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523
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; OTHER INFORMATION:
; US-08-406-855A-5

Query Match          2.7%; Score 34.8; DB 2; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 474 cgcctccagcgcctccagccacggcgcgtggtgcgttatctgcagcgtct 533
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTTCTTCAGCGCT 694

QY 534 ggcacccgtctgtggtgcctctcatctctgtgtcatttgaagagacagtttat 593
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 695 GGGCTCCTTCTACCTGCCTCTGCCATCATCTGCTGATGCTACTGCCGCTACGTTG 754

QY 594 ggaagaagaacccagctggtctcgcgtc 623
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 755 GCCAAGAGGGAGAGCGGGGCTCAAGTC 784

RESULT 15
US-08-722-190-5
; Sequence 5, Application US/08722190
; Patent No. 590128
; GENERAL INFORMATION:
; APPLICANT: Charles Gluchowski, Carlos C. Forray, George
; APPLICANT: Chiu, Theresa A. Brancheke, John M. Wetzel and Paul R. Hartig
; TITLE OF INVENTION: USE OF ALPHA-1C SPECIFIC COMPOUNDS TO
; TITLE OF INVENTION: TREAT BENIGN PROSTATIC HYPERPLASIA
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: COOPER & DUNHAM LLP
; STREET: 1185 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.24
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/722,190
; FILING DATE: 4-APR-1995
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41878-D-PCT/JPW/AGL
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0525
; TELEX:
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHETICAL: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523
; OTHER INFORMATION:
; US-08-722-190-5

Query Match          2.7%; Score 34.8; DB 2; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76;
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	Matches	78;	Conservative	0;	Mismatches	72;	Indels	0;	Gaps	0;
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Db	635	CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCCGGGCTACGTCTCTCTCAGCGCT	594							
QY	534	ggccaccgtcctgctggccctgctcctctctgtgtcatctattgtaagagacagtttat	593							
Db	695	GGGCTCCTTCTACCTGCCTCTGGCCATCATCCTTGGTTCATGTACTGCCGGCTCTACGTGGT	754							
QY	594	ggagaagaaccagctggtctctgcggtc	623							
Db	755	GGCCAAAGAGGGAGAGCCGGGGCTCAAGTC	784							

Search completed: April 11, 2002, 00:22:56
 Job time: 10243 sec

GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: April 11, 2002, 00:22:56 ; Search time 107.54 Seconds
(without alignments)
3588.604 Million cell updates/sec

Title: US-09-380-276A-2
Perfect score: 1704
Sequence: 1 ggggaacgtagaactctccaa.....gaccagagtatacttttc 1704

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA.*
1: /cgn2_6/ptodata/2/ina/5A_COMB.seq.*
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3: /cgn2_6/ptodata/2/ina/6A_COMB.seq.*
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5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq.*
6: /cgn2_6/ptodata/2/ina/backfiles1.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	447.2	26.2	893	4	US-09-386-529-8
2	326.6	19.2	623	4	US-09-386-529-9
3	36.4	2.1	1601	1	US-08-722-001-7
4	36.4	2.1	1987	1	US-08-722-001-26
5	36.4	2.1	1997	1	US-08-722-001-27
6	36.4	2.1	2004	1	US-08-722-001-11
7	36.2	2.1	2485	1	US-08-424-424B-1
8	36.2	2.1	2486	5	PCT-US94-05363A-1
9	36	2.1	4136	4	US-09-103-875-2
10	35.6	2.1	1150	4	US-09-372-934-3
11	35.4	2.1	7218	1	US-08-232-463-14
12	34.8	2.0	1639	1	US-08-334-698-5
13	34.8	2.0	1639	1	US-08-228-932-5
14	34.8	2.0	1639	1	US-08-468-939-5
15	34.8	2.0	1639	2	US-08-406-855A-5
16	34.8	2.0	1639	2	US-08-722-190-5
17	34.8	2.0	1639	3	US-08-244-354-5
18	34.8	2.0	1639	3	US-09-206-899-5
19	34.8	2.0	1639	5	PCT-US95-04203-5
20	34.6	2.0	5962	6	5386025-5
21	33.6	2.0	800	2	US-08-416-603-11
22	33.4	2.0	4360	1	US-08-470-350B-1
23	33	1.9	9472	1	US-08-325-547-9
24	32.8	1.9	2230	1	US-08-200-512-1
25	32.6	1.9	1593	2	US-08-524-828-2
26	32.6	1.9	1593	2	US-08-975-114A-2
27	32.6	1.9	1593	3	US-08-849-281A-2

28	32.6	1.9	2247	2	US-08-524-828-1	Sequence 1, Appli
29	32.6	1.9	2247	1	US-08-975-114A-1	Sequence 1, Appli
C 30	32.6	1.9	3891	1	US-08-480-604A-27	Sequence 27, Appli
C 31	32.6	1.9	3891	2	US-08-405-496A-27	Sequence 27, Appli
C 32	32.6	1.9	3891	4	US-08-915-136-27	Sequence 27, Appli
33	32.4	1.9	1167	1	US-07-960-985-1	Sequence 1, Appli
34	32.4	1.9	1167	2	US-08-496-671-1	Sequence 1, Appli
35	32.4	1.9	1280	4	US-09-096-776B-4	Sequence 4, Appli
36	32.4	1.9	1491	4	US-09-082-092-9	Sequence 9, Appli
37	32.4	1.9	1524	4	US-08-840-767-3	Sequence 3, Appli
38	32.4	1.9	1690	2	US-08-461-812-3	Sequence 3, Appli
39	32.4	1.9	3083	2	US-08-480-994-36	Sequence 36, Appli
40	32.4	1.9	3083	2	US-08-616-844-36	Sequence 36, Appli
41	32.4	1.9	3083	2	US-08-599-654-36	Sequence 36, Appli
42	32.4	1.9	3083	2	US-08-485-573-36	Sequence 36, Appli
43	32.4	1.9	3083	3	US-08-944-868A-36	Sequence 36, Appli
44	32.4	1.9	3083	3	US-08-944-423A-36	Sequence 36, Appli
45	32.4	1.9	3083	3	US-08-925-743-36	Sequence 36, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match	26.2%	Score 447.2;	DB 4;	Length 893;
Best Local Similarity	81.6%	Pred. No. 1.4e-127;		
Matches 529;	Conservative	0;	Mismatches 118;	Indels 1;
Gaps	1;			
Qy	22	aataatacatttgataagaagatggctttaaaagtgtactagaacaagaaaacgt	81	
Db	32	aataaacacgttctggtgagagccatggcactcaagggtctctacctacacagggagcgtgc	91	
Qy	82	tttctactcttttagtattactaggtctattgtcatgtataaagtgtactgtgaacacagag	141	
Db	92	tcttcgtgcattctcttctactccaccctggcatgtataaagtgtgcaaacccggag	151	
Qy	142	actgtagacagcaagaattcaggatcggtctggaacctgtgtccctgtcaaccagtgtg	201	
Db	152	attgcaggcagcaggaattcaagatcgatctggaacctgtgtcctctgtcaacagtgcg	211	
Qy	202	ggccagcagtgagttgtctaagaatgtggtcttcggtatgggaggtacacagtgtg	261	
Db	212	gacctggtgagttgtcccaaggaatgtggtcttcggtatgggaggtacacagtgtg	271	
Qy	262	tgacgtgcggctgcacaggttccagaggtggtgggtcccaagaaatgcaagccctgtc	321	
Db	272	tgccctgcaggccgacccggttcaaggaagactgggtttccagaagtgttaagccatgtg	331	
Qy	322	tggactgcgagttggtgaaccgttttcagaaggcaaatgttccagccaccagtgtgcca	381	
Db	332	cggactgtgcgtggtgaaccgttttcagaaggcgaactgtctcacacaccagtgtgctg	391	
Qy	382	tctgcgggactgttccagagattttatagaagacgaaactgtcggtttcaagaca	441	
Db	392	tctgcgggactgtcctccaggtttttaccoggaagacaaactgtgtgttttcaagaca	451	

; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/232,463
 ; FILING DATE:
 ; CLASSIFICATION: 435
 ; PRIOR APPLICATION DATA:
 ; APPLICATION NUMBER: US/07/935,313
 ; FILING DATE:
 ; APPLICATION NUMBER: EP 91 114 300.6
 ; FILING DATE: 26-AUG-1991
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: BENT, Stephen A.
 ; REGISTRATION NUMBER: 29,768
 ; REFERENCE/DOCKET NUMBER: 30472/114 IMMU
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: (703)836-9300
 ; TELEFAX: (703)683-4109
 ; TELEX: 899149
 ; INFORMATION FOR SEQ ID NO: 14:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 7218 base pairs
 ; TYPE: nucleic acid
 ; STRANDEDNESS: single
 ; TOPOLOGY: linear
 ; IMMEDIATE SOURCE:
 ; CLONE: pTZgpt-F1s
 ;
 ; US-08-232-463-14

Query Match 2.1%; Score 35.4; DB 1; Length 7218;
 Best Local Similarity 6.4%; Pred. No. 2.1;
 Matches 27; Conservative 203; Mismatches 189; Indels 0; Gaps 0;

 QY 11 aactctcaacataataacattatagaagaagatgctttaaagtgctactagaaca 70
 DB 1481 AATTACCTATCTATCACTAGTAAAGAGATAGAGAAATTTGGTACRRRRRRRRRR 1422

 QY 71 agagaaacgttttcaactcttttagtattactaggctattgtcatgtaagtgacttg 130
 DB 1421 RRR 1362

 QY 131 tgaacacagagacttagacacagaaatcaggatcggtctggaactgtgtccctg 190
 DB 1361 RRR 1302

 QY 191 caacagtggtgagcagcatgagttgtctaagaatgtgcttgctatggaggga 250
 DB 1301 RRR 1242

 QY 251 tgacagtggtgacgtgcggtgcacaggttcaaggaggactggggcttcagaaatg 310
 DB 1241 RRR 1182

 QY 311 caagccctgtctgactgcagtggtggaacgctttcagaaggcaaatgttcagccac 370
 DB 1181 RRR 1122

 QY 371 cagtgtgcatctggtgggactgttccaggatttatagaagaaactgtctg 429
 DB 1121 RRR 1063

RESULT 12
 US-08-334-698-5
 ; Sequence 5, Application US/08334598
 ; Patent No. 5556753
 ; GENERAL INFORMATION:
 ; APPLICANT: Jonathan A. Bard et al.
 ; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
 ; TITLE OF INVENTION: Receptors and Uses Thereof
 ; NUMBER OF SEQUENCES: 6
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: COOPER & DUNHAM
 ; STREET: 30 Rockefeller Plaza

; CITY: New York
 ; STATE: New York
 ; COUNTRY: U.S.A.
 ; ZIP: 10112
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.24
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/334,698
 ; FILING DATE:
 ; CLASSIFICATION: 435
 ; PRIOR APPLICATION DATA:
 ; APPLICATION NUMBER: US/07/952,798
 ; FILING DATE:
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: White, John P.
 ; REGISTRATION NUMBER: 28,678
 ; REFERENCE/DOCKET NUMBER: 376901
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: (212) 977-9550
 ; TELEFAX: (212) 664-0525
 ; TELEX: (212) 42523 COOP UI
 ; INFORMATION FOR SEQ ID NO: 5:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 1639 base pairs
 ; TYPE: nucleic acid
 ; STRANDEDNESS: single
 ; TOPOLOGY: unknown
 ; MOLECULE TYPE: DNA (genomic)
 ; HYPOTHETICAL: N
 ; ANTI-SENSE: N
 ; FEATURE:
 ; NAME/KEY: CDS
 ; LOCATION: 126..1523
 ; OTHER INFORMATION:
 ;
 ; US-08-334-698-5

Query Match 2.0%; Score 34.8; DB 1; Length 1639;
 Best Local Similarity 52.0%; Pred. No. 1.3;
 Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

 QY 518 cgcgtccacggcctcagccacgggacacggcgctggtgcgttatctgcagcgtct 577
 DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTGCTCTCTCTCAGCGCT 694

 QY 578 gggccaccgtcctgtgcccctctcatcctctgtgtcatctattgtaagacagattat 637
 DB 695 GGGCTCTTCTACCTGCTCTGCGCATCATCTCTGCTCATCTACTGCGCGCTCTACGTGCT 754

 QY 638 ggagaagaaacccagctggtctctgcggtc 667
 DB 755 GGCCAAGAGGAGAGCGCGGGCTCAAGTC 784

RESULT 13
 US-08-228-932-5
 ; Sequence 5, Application US/08228932
 ; Patent No. 5578611
 ; GENERAL INFORMATION:
 ; APPLICANT: Charles Gluchowski, Carlos C. Forray, George Chiu,
 ; APPLICANT: Theresa A. Branche, John M. Wetzel and Paul R. Hartlg
 ; TITLE OF INVENTION: USE OF ALPHA-1C SPECIFIC COMPOUNDS TO TREAT BENIGN
 ; TITLE OF INVENTION: PROSTATIC HYPERPLASIA
 ; NUMBER OF SEQUENCES: 6
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: COOPER & DUNHAM
 ; STREET: 30 Rockefeller Plaza
 ; CITY: New York
 ; STATE: New York
 ; COUNTRY: U.S.A.

```

1  ZIP: 10112
2
3  COMPUTER READABLE FORM:
4
5  MEDIUM TYPE: Floppy disk
6
7  COMPUTER: IBM PC compatible
8
9  OPERATING SYSTEM: PC-DOS/MS-DOS
10
11  SOFTWARE: Patent In Release #1.24
12
13  CURRENT APPLICATION DATA:
14
15  APPLICATION NUMBER: US/08/228,932
16
17  FILING DATE: 13-APR-1994
18
19  CLASSIFICATION: 514
20
21  ATTORNEY/AGENT INFORMATION:
22
23  NAME: White, John P.
24
25  REGISTRATION NUMBER: 28,678
26
27  REFERENCE/DOCKET NUMBER: 41878-B/JWP/TEP
28
29  TELECOMMUNICATION INFORMATION:
30
31  TELEPHONE: (212) 977-9550
32
33  TELEFAX: (212) 664-0525
34
35  TELEX: (212) 472533 COOP UI
36
37  INFORMATION FOR SEQ ID NO: 5:
38
39  SEQUENCE CHARACTERISTICS:
40
41  LENGTH: 1639 base pairs
42
43  TYPE: nucleic acid
44
45  STRANDEDNESS: single
46
47  TOPOLOGY: unknown
48
49  MOLECULE TYPE: DNA (genomic)
50
51  HYPOTHETICAL: N
52
53  ANTI-SENSE: N
54
55  FEATURE:
56
57  NAME/KEY: CDS
58
59  LOCATION: 126..1523
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61  OTHER INFORMATION:
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63  US-08-228-932-5

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	Query Match	2.0%	Score 34.8;	DB 1;	Length 1639;
	Best Local Similarity	52.0%;	Pred. No. 1.3;		
	Matches	78;	Conservative	0;	Mismatches 72; Indels 0; Gaps 0;
Qy	518	cgggtccacggctccagccacgggacagggcctggctgcgcttatctcagcgcctct	577		
Db	635	CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCCGGGCTACGTGCTCTTCTCAGCGCT	694		
Qy	578	ggccacgcctcgtgtggccctgctcctctgtgtcatctatgtgaagacacagtttat	637		
Db	695	GGGCTCCTTCTACTCTGCTCTGGCCATCATCTCGTCTATGTACGTGCCCGCTCAGTGCT	754		
Qy	638	ggagaagaaccccgctggtctctgcggctc	667		
Db	755	GGCCAAGAGGGAGAGCCGGGGCTTCAGTC	784		

```

RESULT 14
US-08-468-939-5
; Sequence 5, Application US/08468939
; Patent No. 5714381
; GENERAL INFORMATION:
; APPLICANT: Jonathan A. Bard et al.
; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
; TITLE OF INVENTION: Receptors and Uses Thereof
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: COOPER & DUNHAM LLP
; STREET: 1185 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.24
; CURRENT APPLICATION DATA:

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; APPLICATION NUMBER: US/08/468,939
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41337-1b
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0526
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHEetical: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523
; OTHER INFORMATION:
;
US-08-468-939-5

Query Match          2.0%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 1.3;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 518 cgcgctcacggcctccagccacggggacagggcgctggctgcgcttatctgcagcgctct 577
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGACCGCGGCTACTGTCTTCTTCAGCGCT 694
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 578 ggcacccgctcgtcgtggccctgctcctctctgtgctatctattgtaagagacagtttat 637
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 695 GGGCTCCTTCTACCTGGCTCTGGCCATCACTCTGGTCTATGTACTGCCCGCTCTACGTGGT 754
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 638 ggagaagaacccagctggtctctgcgctc 667
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 755 GGCCACAGGGGAGAGACCGGGGCGCTCAAGTC 784
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RESULT 15
US-08-406-855A-5
Sequence 5, Application US/08406855A
Patent No. 5861309
GENERAL INFORMATION:
APPLICANT: Jonathan A. Bard et al.
TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
Receptors and Uses Thereof
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/406,855A
FILING DATE: 21-Aug-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41337-A-PCT-US/JPW/KDB
TELECOMMUNICATION INFORMATION:

GenCore version 4.5
Copyright (c) 1993 - 2000 Compuqen Ltd.

OM protein - protein search, using sw model

Run on: April 10, 2002, 17:14:57 ; Search time 21.63 Seconds
(without alignments)
433.836 Million cell updates/sec

Title: US-09-380-276A-4
Perfect score: 2255
Sequence: 1 MALKVLLQEKTFTLLVLL.....LDQESGAIHPATOTSLOEA 417

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 212252 seqs, 22503292 residues

Total number of hits satisfying chosen parameters: 212252

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query %		DB	ID	Description
		Match	Length			
1	869	38.5	210	4	US-09-286-529-3	Sequence 3, Appli
2	714.5	31.7	151	4	US-09-286-529-4	Sequence 4, Appli
3		7.1	438	1	US-08-097-827-11	Sequence 11, Appl
4	159	7.1	438	1	US-08-494-574-11	Sequence 11, Appl
5	150	6.7	206	1	US-08-097-827-7	Sequence 7, Appli
6	150	6.7	206	1	US-08-494-574-7	Sequence 7, Appli
7	145	6.4	205	3	US-08-974-022-51	Sequence 51, Appl
8	145	6.4	205	4	US-08-795-445A-51	Sequence 51, Appl
9	145	6.4	205	4	US-08-795-447A-51	Sequence 51, Appl
10	145	6.4	205	4	US-08-974-186-51	Sequence 51, Appl
11	145	6.4	205	4	US-08-795-446B-51	Sequence 51, Appl
12	144	6.4	1104	2	US-08-327-832-5	Sequence 5, Appli
13	144	6.4	1104	2	US-08-828-584-5	Sequence 5, Appli
14	136	6.0	625	3	US-08-996-139-15	Sequence 15, Appl
15	136	6.0	625	4	US-08-995-659-15	Sequence 15, Appl
16	136	6.0	625	4	US-09-215-649A-15	Sequence 15, Appl
17	134.5	6.0	415	4	US-09-006-353A-6	Sequence 6, Appli
18	134	5.9	186	1	US-08-089-458B-6	Sequence 6, Appli
19	134	5.9	307	4	US-08-804-166-4	Sequence 4, Appli
20	134	5.9	307	4	US-08-910-991-4	Sequence 4, Appli
21	133.5	5.9	2050	2	US-08-347-594A-2	Sequence 2, Appli
22	132.5	5.9	197	2	US-08-505-606-1	Sequence 1, Appli
23	132	5.9	276	4	US-09-041-886-27	Sequence 27, Appl
24	132	5.9	277	4	US-09-042-785A-10	Sequence 10, Appl
25	132	5.9	277	4	US-09-006-353A-10	Sequence 10, Appl
26	131	5.8	139	2	US-08-219-237B-8	Sequence 8, Appli
27	131	5.8	176	4	US-09-411-722-1	Sequence 1, Appli

ALIGNMENTS

RESULT 1

```

US-09-286-529-3
; Sequence 3, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AN
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0.
; SEQ ID NO 3
; LENGTH: 210
; TYPE: PRT
; ORGANISM: human
; US-09-286-529-3

```

Query Match	38.5%	Score 869;	DB 4;	Length 210;
Best Local Similarity	83.7%;	Pred. No. 3.9e-73;		
Matches 154:	Conservative	11: Mismatches	19: Indels	Gaps 0:

[illegible]

Ov 181 VILLA 184

Db 187 VI.LA 184

RESULTS

RESULT
 US-09-286-529-4
 : Sequence 4, Application US/09286529
 : Patent No. 6297367
 : GENERAL INFORMATION:
 : APPLICANT: Catherine Tribouley
 : TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
 : FILE REFERENCE: 1408.003/200130.439C1
 : CURRENT APPLICATION NUMBER: US/09/286,529
 : CURRENT FILING DATE: 1999-04-05

NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 151
; TYPE: PRT
; ORGANISM: human
US-09-286-529-4

Query Match 31.7%; Score 714.5; DB 4; Length 151;
Best Local Similarity 82.0%; Pred. No. 5.5e-59;
Matches 123; Conservative 9; Mismatches 17; Indels 1; Gaps 1;
QY 1 MALKVLEQKEFFLLVLLGLSKVTCETGDC-RQEFDRSGNCVPCNOCGPGMELS 59
Db 1 MALKVLP LHRVTF AAILLLHACKVSCETGDCSRQEFKDRSGNVLCKCGPGMELS 60
QY 60 KECFGYGEDACVCTRLHREFKEDMGFKCKPCCLDCAVVNRFOKANCATSATSDAIGDCCLP 119
Db 61 KECFGYGEDACVCPCHRRFKEDMGFKCKPCCLDCALVNRFRANCSTSDAVGDCCLP 120
QY 120 GFYRKTGLVGFODMECVCGDPPPPVPEHC 149
Db 121 GFYRKTGLVGFODMECVCGDPPPPVPEHC 150

RESULT 3
US-08-097-827-11
; Sequence 11, Application US/08097827
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; Goodwin, Ray
; Fanslow, William
; Gayle, Richard
; TITLE OF INVENTION: Novel Cytokine which is a Ligand for
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-Jul-1993
; CLASSIFICATION: <unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 438 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 11:
US-08-097-827-11

Query Match 7.1%; Score 159; DB 1; Length 438;
Best Local Similarity 27.5%; Pred. No. 9.9e-07;
Matches 69; Conservative 31; Mismatches 93; Indels 58; Gaps 18;

QY 15 TLVLVLGLYLSCKVTCETGDCRQEQEFDRSGN-CVPCNOCGPGMELSKECGFGYGEDAQC 73
Db 9 TALLLLG-LTLGVTRRLNCVKHTY--PSGKHC--CRECQPGHGMVNR--DHTRTLTCH 61
QY 74 TCRHLRHKEDMGFKCKPCCLDCAVVNRFO-KANCATSATSDAIGDCCLPGLPYRKTGLVGFOD 132
Db 62 PCETGFYNEAVNYDTCKQCTQCNHRSGLKQNCPTQDTVC-RCRPGTQPR-----QD 114
QY 133 -----MECVCGDPPPPVPEP-----HCASKVNLVKLTASTASSPRDTALAAVIC---SALA 179
Db 115 SGYKLGVDVCP--PPGHFSPGNQACKPWTNCTLSGKOTRHPASDSDAV-CEDRSLLA 171
QY 180 TVLLALLLILCVYCKRQFMEKPPSW---SLRSODIQYNGSELSCLDRLPOLHEVAHRACQ 236
Db 172 TLL-----WETQRTFPTTVQSTVWPTSELP--STPTLVE--PRSC--- 211
QY 237 CRDSVQTCGP 247
Db 212 ---DKTHTCPP 219
RESULT 4
US-08-494-574-11
; Sequence 11, Application US/08494574
; Patent No. 5783665
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; APPLICANT: Goodwin, Ray
; APPLICANT: Fanslow, William
; APPLICANT: Gayle, Richard
; TITLE OF INVENTION: No. 5783665el Cytokine which is a Ligand for
; TITLE OF INVENTION: OX40
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/494,574
; FILING DATE: 22-JUN-1995
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-JUL-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 438 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-494-574-11

Query Match 7.1%; Score 159; DB 1; Length 438;
Best Local Similarity 27.5%; Pred. No. 9.9e-07;
Matches 69; Conservative 31; Mismatches 93; Indels 58; Gaps 18;
QY 15 TLVLVLGLYLSCKVTCETGDCRQEQEFDRSGN-CVPCNOCGPGMELSKECGFGYGEDAQC 73
Db 9 TALLLLG-LTLGVTRRLNCVKHTY--PSGKHC--CRECQPGHGMVNR--DHTRTLTCH 61

Db 9 TALLLLG-LTLGVTAARRLNCVKHTY--PSGHKC--CRECOPGHGMVNR--DHTRDTLCH 61
QY 74 TCRHLRFKEDWGQKCKPCLDCAVNRFO-KANCATSDAICGDCPLPGFYRKTCLVGFOD 132
Db 62 PCETGYNEAVNYDTCQCTQCNHRSGSELKQNCPTQDTVC-RCRPGTQPR-----QD 114
QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIATASSPRDTALAAVIC---SALA 179
Db 115 SGYKLGVDVCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171
QY 180 TVLLALLILCVYKQFMEKKPSW---SLRSODIOYNGSELSCLDRLPOLHEYAHRACCO 236
Db 172 TLL-----WETQRTPTRTTVQSTVWPRTSSELP--STPTLVE--PRSC-- 211
QY 237 CRDVSQTCGP 247
Db 212 ---DKTHTCPP 219

RESULT 5

US-08-097-827-7
; Sequence 7, Application US/08097827
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; Goodwin, Ray
; Fanslow, William
; Gayle, Richard
; TITLE OF INVENTION: Novel Cytokine Which is a Ligand for
; OX40
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-Jul-1993
; CLASSIFICATION: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 206 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-08-097-827-7

Query Match 6.7%; Score 150; DB 1; Length 206;
Best Local Similarity 30.1%; Pred. No. 2.4e-06;
Matches 55; Conservative 21; Mismatches 75; Indels 32; Gaps 13;

QY 15 TLVLVLGLYSCKVTCGTGCRQOEFRDRSGN-CVPCNOCGPGMELSKGCGFGYGEDAQCV 73
Db 9 TALLLLG-LTLGVTAARRLNCVKHTY--PSGHKC--CRECOPGHGMVNR--DHTRDTLCH 61
QY 74 TCRHLRFKEDWGQKCKPCLDCAVNRFO-KANCATSDAICGDCPLPGFYRKTCLVGFOD 132
Db 62 PCETGYNEAVNYDTCQCTQCNHRSGSELKQNCPTQDTVC-RCRPGTQPR-----QD 114

QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIATASSPRDTALAAVIC---SALA 179
Db 115 SGYKLGVDVCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171
QY 180 TVL 182
Db 172 TLL 174

RESULT 6

US-08-494-574-7
; Sequence 7, Application US/08494574
; Patent No. 5783665
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; APPLICANT: Goodwin, Ray
; APPLICANT: Fanslow, William
; APPLICANT: Gayle, Richard
; TITLE OF INVENTION: No. 5783665el Cytokine Which is a Ligand for
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/494,574
; FILING DATE: 22-JUN-1995
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-JUL-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 206 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-494-574-7

Query Match 6.7%; Score 150; DB 1; Length 206;
Best Local Similarity 30.1%; Pred. No. 2.4e-06;
Matches 55; Conservative 21; Mismatches 75; Indels 32; Gaps 13;

QY 15 TLVLVLGLYSCKVTCGTGCRQOEFRDRSGN-CVPCNOCGPGMELSKGCGFGYGEDAQCV 73
Db 9 TALLLLG-LTLGVTAARRLNCVKHTY--PSGHKC--CRECOPGHGMVNR--DHTRDTLCH 61
QY 74 TCRHLRFKEDWGQKCKPCLDCAVNRFO-KANCATSDAICGDCPLPGFYRKTCLVGFOD 132
Db 62 PCETGYNEAVNYDTCQCTQCNHRSGSELKQNCPTQDTVC-RCRPGTQPR-----QD 114
QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIATASSPRDTALAAVIC---SALA 179
Db 115 SGYKLGVDVCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171
QY 180 TVL 182

Db 172 TLL 174
!:

RESULT 7

US-08-974-022-51
; Sequence 51, Application US/08974022
; Patent No. 6015938
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/974,022
; FILING DATE: 12-DEC-1995
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-974-022-51

Query Match 6.4%; Score 145; DB 3; Length 205;
Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QKTFFTLLVLLGLSKVTCETGDCRQOEFRDR--SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QQPTAFLLGLSLGVTVKLNC-----VKDTPSGHKC--CRECQPGHGMVSR--D 52

QY 66 YGEDAQCVCRLHRFKEDMGFKCKPCLDCAVNVNRQ--KANCSTASDAICGDCPLPGFYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKQCTOCNHRSGSELKQNCPTPTDVTVC-QCRPGCTQPR 111

QY 125 TKLVGFQDMCEVCPGDPDPPEY-----HCASKYNLV-----KIASTASSPRDTALAAVIC- 175
Db 112 QDSSHLKLGVDVCP--PPGHFSPGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164

QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 8
US-08-795-445A-51
; Sequence 51, Application US/08795445A
; Patent No. 6284485
; GENERAL INFORMATION:

; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,445A
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-795-445A-51

Query Match 6.4%; Score 145; DB 4; Length 205;
Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QKTFFTLLVLLGLSKVTCETGDCRQOEFRDR--SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QQPTAFLLGLSLGVTVKLNC-----VKDTPSGHKC--CRECQPGHGMVSR--D 52

QY 66 YGEDAQCVCRLHRFKEDMGFKCKPCLDCAVNVNRQ--KANCSTASDAICGDCPLPGFYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKQCTOCNHRSGSELKQNCPTPTDVTVC-QCRPGCTQPR 111

QY 125 TKLVGFQDMCEVCPGDPDPPEY-----HCASKYNLV-----KIASTASSPRDTALAAVIC- 175
Db 112 QDSSHLKLGVDVCP--PPGHFSPGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164

QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 9
US-08-795-447A-51
; Sequence 51, Application US/08795447A
; Patent No. 6284728
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: Osteoprotegerin
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: One Amgen Center Drive

LENGTH: 205 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-795-446B-51

Query Match
Best Local Similarity 6.4%; Score 145; DB 4; Length 205;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QKTEFTLLVLGLYSCKVTCTGCRQOEFRDR--SGN-CVPCNQCGMELSKCEGFG 65
DB 6 QQTAFLLLGLSLGVTYKLNLC-----VKDTYPSGHKC--CREQPGHGMVSRD--D 52
QY 66 YGEDACVTCRLHRRKEDMGFKKPCLDCAVNNRFQ-KANGSATSDAICGDLPGFYRK 124
DB 53 HTRDTVCHPCPEGFYNEAVNYDTCKOCTQCNHRSGSELKQNTPTEDTVC-QCRPGTQPR 111
QY 125 TKLVGFQDMCEVPCGDPPPPEP---HCASKVNLV---KIATASSPRDTALAAVIC- 175
DB 112 QSSSHKLVDCVPC--PPGHFSGNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164
QY 176 --SALATVL 182
DB 165 DRSLATL 173

RESULT 12
US-08-327-832-5
; Sequence 5, Application US/08327832
; Patent No. 5840832

GENERAL INFORMATION:
APPLICANT: Ono, Santa J.
TITLE OF INVENTION: Transcription Factor Regulating MHC
TITLE OF INVENTION: Expression, cDNA and Genomic Clones Encoding Same and
TITLE OF INVENTION: Retroviral Expression Contracts Thereof
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner, Birch, McKie & Beckett
STREET: 1001 G Street, N.W.
CITY: Washington, D.C.
STATE: District of Columbia
COUNTRY: U.S.A.
ZIP: 20001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/327,832
FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Posorske, Laurence H.
REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 1107,46362
TELECOMMUNICATION INFORMATION:
TELEPHONE: 20-2 508-9153
TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1104 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-327-832-5

Query Match 6.4%; Score 144; DB 2; Length 1104;

Best Local Similarity 18.8%; Pred. No. 8.8e-05;
Matches 110; Conservative 59; Mismatches 181; Indels 236; Gaps 27;
QY 24 SCKVTCGTCRQOEFRDRSGNCVPCNQCGMELSKCEG-----FGYGEDAQC----- 72
DB 440 SCNLLCHPG-----PCPPCPAPMTKTCEGTRHTVRCQGVSVHCSNPC 484
QY 73 ---VTCRLHRRKEDMGFKKPCLDCAVNNRFQKANGSATS-DAICGDLCPGFYRKTKLV 128
DB 485 ENILNCGQHQAEHLCHGGCQPCQ--IILN--OVYCGSTSRDVLJGTDV-----GKSD 534
QY 129 GFQDMCE-----VPCGD-----PPPP-----PYEPHC--ASKYNLVKIASTASS 164
DB 535 GFGDFSLCTGCKDLKGNHCSQVCHPQPCQCPRLPOLVRCCPGQGPLSQLLLELSS 594
QY 165 PRDTALAIV-----TC-SALATVLLALLILC-----VYICKRQFMEKK- 201
DB 595 SRKTCMDPVPCGKVGKPLPGSLDFIHTCEKLCHEGDCGPVSRITSVISCRCSEFTKEL 654
QY 202 PMSLRSQDI-----QYNGSELSCLDPRQ-----LH-----EYA 230
DB 655 PCTSLKSEDATEMCDKRCNKKRLCGRHKCNELCCVDKEHKCPLNCGRLKRCGLHRCPEPC 714
QY 231 HRACCO-CRRDSVQT-----CGPVRLPLSMCC-----EE 258
DB 715 HRGNCQTCWQASFDLTCGASVIYPPVPCGTRPPECTQTCARVHECDHPVHSGHSEE 774
QY 259 ACS-----PMPATLGGGVHISAASL----- 277
DB 775 KCPCTCTLTQKCMGKHFRSNIPCHLVDISGLPCSATLPCGMHMKCQRLCHKGECLVDE 834
QY 278 -----QAR-----NAGPAGEWPTFFGSLTQSI 300
DB 835 PCKQPCCTTPRADCCHPCMAPCHTSSPCPVACKAKAVELOCEGCRKRMVICSEASTYOR 894
QY 301 CGEFSDAWPLMQNMGDMISFCDSPYELTGEDIHSLNPELESSTSLDSSNQDLVGGAV 360
DB 895 IAAISMASKITDMLGGS-----VEISKLTIKKEVHQARLECEDECSALERKKR--LAEAF 948
QY 361 PVQSHSENFATDLSRYNNTLVESASTODALTMRSQLDOESGAI 406
DB 949 HISEDPDFNIRSSGSKFSDSLKEDA--RKDLKFVSDVEKEMETLV 992
RESULT 13
US-08-828-584-5
; Sequence 5, Application US/08828584
; Patent No. 5908762
GENERAL INFORMATION:
APPLICANT: Ono, Santa J.
TITLE OF INVENTION: Transcription Factor Regulating MHC
TITLE OF INVENTION: Expression, cDNA and Genomic Clones Encoding Same and
TITLE OF INVENTION: Retroviral Expression Contracts Thereof
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner, Birch, McKie & Beckett
STREET: 1001 G Street, N.W.
CITY: Washington, D.C.
STATE: District of Columbia
COUNTRY: U.S.A.
ZIP: 20001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,584
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Posorske, Laurence H.

REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 1107.46362
TELECOMMUNICATION INFORMATION:
TELEPHONE: 20-2 508-9153
TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1104 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-828-584-5

Query Match 6.4%; Score 144; DB 2; Length 1104;
Best Local Similarity 18.8%; Pred. No. 8.8e-05;
Matches 110; Conservative 59; Mismatches 181; Indels 236; Gaps 27;

QY 24 SKKVTCTGDCRQEQFRDRSGNVCNQCQPGOMELSKECG-----FGYGEDAOC----- 72
DB 440 SCNLLCHPG-----PCPPCPAFMTKTCCGTRTHRVRCGQAVSVHCSNPC 484
QY 73 ---VTCRLHRRFKEDWFOKRCPLDCAVVNRFOKANCATS-DAICGDCPLGCFYRKTILV 128
DB 485 ENILNCGQHQAELCHGGCCQPCQ--IILN--QVCYCGSTRDVLGCTDV-----GKSD 534
QY 129 GFQDMEC-----VPCGD-----PYEPHC--ASKVNLVKIASTASS 164
DB 535 GDFSCSLCTGCKDLKGNHTCSQVCHPOCQCPRLPOLVRCPCGQTPLSOLLELGS 594
QY 165 PRDTALAAV-----TC-SALATVILALLILC-----VIYKRRQFMKK- 201
DB 595 SRKTCMDPVPCGKVGKPLPCGSLDFIHTCEKLGHEGDCGPVSRVTSVISCRCSPRTKEL 654
QY 202 PWSLSRSODI-----QYNGSELSCLDLDPQ-----LH---EVA 230
DB 655 PCTSLKSEDATEMCKDKRNNKBLGRHCKNCCVDKHEKPLNCGRLRGLHRCCEPC 714
QY 231 HRACQ-CRRDSVQT-----CGPVRLLPSMCC-----EE 258
DB 715 HRGNCOTCWOASFDELTHCHGASVIYPPVPCGTRPCTQTCARVHCECDHPYHSHSEE 774
QY 259 ACS-----PNPATLGGCVHSAASL----- 277
DB 775 KCPCTFLTQKCMGKHEFRSNIPCHLVDISGLPCSATPLCGMHKCORLCHKHCECLVDE 834
QY 278 -----QAR-----NAGPAGEMVPTFFGSILTQSI 300
DB 835 PCKQCTTPRADCGHPCMAPHTSSPCPVYACKAKVLOCEGCRKRWICSEASTYQR 894
QY 301 CGEFSADWPLMONGDNISFCDSPYELTGEDIHSLNPELESSTSLDSNSSQDLVGGAV 360
DB 895 IAAISMASKITDMLGGS-----VEISKLTKEVHOARLECEDECSALERKKR--LAEAF 948
QY 361 PVQSHENFTATDLSRYNNTLVESASTQDALTMRSQDOESCAII 406
DB 949 HISEDSDPENIRSSGSKFSDSLKEDA--RKDLKFVSDVEKEMETLV 992

RESULT 14

US-08-996-139-15
Sequence 15, Application US/08996139
Patent No. 6017729
GENERAL INFORMATION:
APPLICANT: Anderson, Dirk M.
APPLICANT: Galibert, Laurent
APPLICANT: Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immunex Corporation, Law Department
STREET: 51 University Street
CITY: Seattle

STATE: WA
COUNTRY: USA
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple operating System 7.5.5
SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/996,139
FILING DATE: 22 DECEMBER 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 60/064,671
FILING DATE: 14 OCTOBER 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2851-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 625 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-996-139-15

Query Match 6.0%; Score 136; DB 3; Length 625;
Best Local Similarity 22.2%; Pred. No. 0.00022;
Matches 102; Conservative 51; Mismatches 171; Indels 136; Gaps 27;

QY 16 LVLVLGLYSCKVTCE-TGDCRQEQFRDRSGNVCNQCQPGMELSKECGFGYGEDAQCVT 74
DB 18 LCVLL--VPLQVTLQVTPCTQERHYEHLGRG--CSRCEPGYLSKSC--TPTSDSVCLP 71
QY 75 CRLHREKEDWGFQ-KC---KPC-LDCAVV-----NRFOKANCAT-----SDAIC--- 114
DB 72 CGPDEYLDTWNEEDKLLHKKVCDACKALVAVDPGNHTAPRCACCTAGYHWNDSCECCRN 131
QY 115 GDLCPGFYRKTILVGFQDMECVPC-----GDPPPPY-----BPH----- 148
DB 132 TECAPGFGAHPQLNKNKDTVCTPCLLGFSDVFSSTDKCKPWTNCTLLGKLEAHQGTES 191
QY 149 ---CASKVNLVKIASTASSPRDTALAAVICSALATVLLALLILCVYKRRQFMKKPSWS 205
DB 192 DVCSSMTLRRPKEAQAYLPSLI--VLLLFISVVVAAIIFGVYIRKGGKALTANLWN 249
QY 206 -LRSQDIOYNGSELSCLDRLPQLHEYAHRAAC---QCRRDSVOTCGPVRLLPSMCCCEACS 261
DB 250 WYNDACSSLSGNKESGGDR-----CAGSHSTSSQOEVCCEGILL---MTRKKMV 296
QY 262 PNPATLGGCVHSAAS-----LOARNAGPAGEMVPT-----FFGS 295
DB 297 PEDGAGVCGPVCAAGGPAEVRDSRTFTLVSEVETQGLSRKIPTDEYTRDPSQSTGS 356
QY 296 LTQSTCGEESDAWPLMQNPM---GGDNISFC-----DSPELT 330
DB 357 LL--LIQOGSKSIPFPQEPLEVGENDSLSQCFGTGTSTVDSGCDFTPEPPTSDSMP--V 412
QY 331 GEDIHSLNPELESSTSL-----DSNSSQDLVG--GAVPVQSH 365
DB 413 SPEKH-LIKEIEGDSCLPWWVSSNSTDGYTSGNTPGEDH 451

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GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run On: April 11, 2002, 00:23:10 ; Search time 107.54 Seconds
(without alignments)
2672.499 Million cell updates/sec

Title: US-09-380-276A-5
Perfect score: 1269
Sequence: 1 atggcttaaaagtctact.....ggcagcgactgggttcctcg 1269

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	435.4	34.3	893	4	US-09-286-529-8
2	314	24.7	623	4	US-09-286-529-9
3	36.4	2.9	1501	1	US-08-722-001-7
4	36.4	2.9	1987	1	US-08-722-001-26
5	36.4	2.9	1997	1	US-08-722-001-27
6	36.4	2.9	2004	1	US-08-722-001-11
7	36.2	2.9	2485	1	US-08-424-424B-1
8	36.2	2.9	2486	5	PCT-US94-05363A-1
9	36	2.8	4136	4	US-09-103-875-2
10	35.6	2.8	1130	4	US-09-372-934-3
11	34.8	2.7	1639	1	US-08-334-698-5
12	34.8	2.7	1639	1	US-08-228-932-5
13	34.8	2.7	1639	1	US-08-468-935-5
14	34.8	2.7	1639	2	US-08-406-855A-5
15	34.8	2.7	1639	2	US-08-722-190-5
16	34.8	2.7	1639	3	US-08-244-354-5
17	34.8	2.7	1639	3	US-09-206-899-5
18	34.8	2.7	1639	5	PCT-US95-04203-5
19	34.6	2.7	5962	6	US95025-5
20	34.6	2.7	7218	1	US-08-232-463-14
21	33.6	2.6	800	2	US-08-416-603-11
22	33.4	2.6	4360	1	US-08-470-350B-1
23	33	2.6	9472	1	US-08-325-547-9
24	32.8	2.6	2230	1	US-08-200-512-1
25	32.6	2.6	1593	2	US-08-524-828-2
26	32.6	2.6	1593	2	US-08-975-114A-2
27	32.6	2.6	1593	3	US-08-849-281A-2

28	32.6	2.6	2247	2	US-08-524-828-1	Sequence 1, Appli
29	32.6	2.6	2247	2	US-08-975-114A-1	Sequence 1, Appli
c 30	32.6	2.6	3891	1	US-08-480-604A-27	Sequence 27, Appl
c 31	32.6	2.6	3891	2	US-08-405-496A-27	Sequence 27, Appl
c 32	32.6	2.6	3891	4	US-08-915-136-27	Sequence 27, Appl
33	32.4	2.6	1167	1	US-07-960-985-1	Sequence 1, Appli
34	32.4	2.6	1167	2	US-08-496-671-1	Sequence 1, Appli
35	32.4	2.6	1280	4	US-09-096-776B-4	Sequence 4, Appli
36	32.4	2.6	1491	4	US-09-082-092-9	Sequence 9, Appli
37	32.4	2.6	1524	4	US-08-840-767-3	Sequence 3, Appli
c 38	32.4	2.6	1690	2	US-08-461-812-3	Sequence 3, Appli
39	32.4	2.6	3083	2	US-08-480-994-36	Sequence 36, Appl
40	32.4	2.6	3083	2	US-08-616-844-36	Sequence 36, Appl
41	32.4	2.6	3083	2	US-08-599-654-36	Sequence 36, Appl
42	32.4	2.6	3083	2	US-08-485-573-36	Sequence 36, Appl
43	32.4	2.6	3083	3	US-08-944-868A-36	Sequence 36, Appl
44	32.4	2.6	3083	3	US-08-944-423A-36	Sequence 36, Appl
45	32.4	2.6	3083	3	US-08-925-743-36	Sequence 36, Appl

ALIGNMENTS

RESULT 1

US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catheline Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match 34.3%; Score 435.4; DB 4; Length 893;
Best Local Similarity 82.1%; Pred. No. 1.le-125;
Matches 513; Conservative 0; Mismatches 111; Indels 1; Gaps 1;

Qy	1	atggcttttaaaagtctactagaacaagaaacagtttttctactcttttagtattacta	60
Db	55	atgggactcaaggctctactctctacacaggagcgtgtctctcgtccatctcttcta	114
Qy	61	ggctattgtcatgtataaagtgttccctgcacaccagtgtggccaggcaggaattcaag	120
Db	115	ctccacctggcagtataaagtgtgtgcgaacacggagattgcaggcagcaggaattcaag	174
Qy	121	gacggtctggaacagtgttccctgcacaccagtgtggccaggcaggaattgtcttaag	180
Db	175	gacgctatctggaaactgtctctctgcacacagtgcgaacacggagattgcaggcaggaattcaag	234
Qy	181	gaatgtggtctcgctatgggagagatgcacagtgtgtgcggtgcgcagcaggttc	240
Db	235	gaatgtggtctcgctatgggagagatgcacagtgtgtgcggtgcgcagcaggttc	294
Qy	241	aaggagactggggtctccagaataatcaagccctgtctgactgcgcagtggtgaaaccgc	300
Db	295	aaggagactggggtctccagaataatcaagccctgtctgactgcgcagtggtgaaaccgc	354
Qy	301	tttcagaaggaattgttcagccaccagtgtatgcctatcgcgggactgcttgcaggga	360
Db	355	tttcagagggccaactgctcacacaccagtgtatgcctatcgcgggactgcttgcaggga	414
Qy	361	ttttatagaagaacgaaactgtcggctttcaagacatgagtggtgcttcttgagagac	420
Db	415	ttttaccggaagaccacaaactggtgtgtttcaagacatgagtggtgcttctcgagagac	474

REFERENCE/DOCKET NUMBER: 325800-118
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2486 BASE PAIRS
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
TOPOLOGY: LINEAR
MOLECULE TYPE: CDNA
PCT-US94-05363A-1

Query Match 2.9%; Score 36.2; DB 5; Length 2486;
Best Local Similarity 56.2%; Pred. No. 0.36;
Matches 68; Conservative 0; Mismatches 53; Indels 0; Gaps 0;

QY 453 caaggtcaacctgtgaagatgcgtccacggcctccagccacgggacagcgcgctggc 512
DB 2024 CAGCATCATCCAGCTGGGGGTACGCCCGCGCTACAGCGCTGGATCAAGGAGGCG 2083
QY 513 tgccttatctgcagcgctctggccacgcgtcctgctgcccctgctcctctgtgtcat 572
DB 2084 TCCCGAGGCTACCTGTATTTCCTCCCACTGGCCCATGGCACTCCTGTGATCAACCTCATCGT 2143
QY 573 c 573
DB 2144 C 2144

RESULT 9

US-09-103-875-2/c
Sequence 2, Application US/09103875A
Patent No. 6221849

GENERAL INFORMATION:

APPLICANT: Szyf, Moshe
APPLICANT: Bigey, Pascal
APPLICANT: Ramchandani, Shyam
TITLE OF INVENTION: DNA METHYLTRANSFERASE GENOMIC SEQUENCES AND ANTISENSE
TITLE OF INVENTION: OLIGONUCLEOTIDES
FILE REFERENCE: 106101.194
CURRENT APPLICATION NUMBER: US/09/103,875A
CURRENT FILING DATE: 1998-06-24
EARLIER APPLICATION NUMBER: 60/069,865
EARLIER FILING DATE: 1997-12-17
EARLIER APPLICATION NUMBER: 08/866,340
EARLIER FILING DATE: 1997-05-30
NUMBER OF SEQ ID NOS: 138
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2

LENGTH: 4136

TYPE: DNA

ORGANISM: Homo sapiens

US-09-103-875-2

Query Match 2.8%; Score 36; DB 4; Length 4136;
Best Local Similarity 49.5%; Pred. No. 0.56;
Matches 93; Conservative 0; Mismatches 95; Indels 0; Gaps 0;

QY 689 cccacagagcctgctgcaagtgcgcctgactcagtcagtcagcctgagggcgcgctg 748
DB 3917 CCACCCACGCGCCCTGCCTGTCTCCCTGAGTCCGTGTTCCCGCCATGTACTACCGCC 3858
QY 749 tcttccatccatgtgctgtgagggcctcgagcccaaccgcgactctgtgttg 808
DB 3857 TCGGACATCTGTGGGACGACGAGATGGCCGACGCCAGTGTGGCACCCTGGGG 3798
QY 809 ggggtcattctgcagccagctcttcagggaagaacgagccagccgagggagatggtgc 868
DB 3797 CGGTACGCGCGCATCTCGGAGGCTTTCAGACAGCGCGCGGCGGACGAGCGGCC 3738

QY 869 cgactttc 876
DB 3737 GGCITTTT 3730

RESULT 10

US-09-372-934-3
Sequence 3, Application US/09372934
Patent No. 6248579

GENERAL INFORMATION:

APPLICANT: Stutzman-Engwall, Kim J.
APPLICANT: McArthur, Hamish
APPLICANT: Katoh, Yoshihiro
TITLE OF INVENTION: STREPTOMYCES AVERMITILIS GENE DIRECTING THE RATIO OF
FILE REFERENCE: PC10649
CURRENT APPLICATION NUMBER: US/09/372,934
CURRENT FILING DATE: 1999-08-12
EARLIER APPLICATION NUMBER: 60/074,636
EARLIER FILING DATE: 1998-02-13
EARLIER APPLICATION NUMBER: PCT/IB99/00130
EARLIER FILING DATE: 1999-01-25
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 1150
TYPE: DNA
ORGANISM: Streptomyces hygroscopicus
NAME/KEY: CDS
LOCATION: (58)..(990)
US-09-372-934-3

Query Match 2.8%; Score 35.6; DB 4; Length 1150;
Best Local Similarity 57.0%; Pred. No. 0.35;
Matches 65; Conservative 0; Mismatches 49; Indels 0; Gaps 0;

QY 448 gccagcaagtcacactgtgtagatcgctccacggcctccagccacgggacagcgcg 507
DB 142 gccagacggcgctacgcgtatcgagaagcgctcccgcccgccagggcggtgggactcgag 201
QY 508 ctgggtgcggttatctgtgcagcgctctggccacgctctgctgctgctgctgctc 561
DB 202 cggatgcgcggtgtgtgctgctgctgctgctgctgctgctgctgctgctgctc 255

RESULT 11

US-08-334-698-5
Sequence 5, Application US/08334698
Patent No. 5556753

GENERAL INFORMATION:

APPLICANT: Jonathan A. Bard et al.
TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
TITLE OF INVENTION: Receptors and Uses Thereof
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM
STREET: 30 Rockefeller Plaza
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10112

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/334,698
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/07/952,798
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: White, John P. 28,678
REGISTRATION NUMBER: 376901
REFERENCE/DOCKET NUMBER: 376901
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEFAX: (212) 664-0525
TELEX: (212) 422523 COOP UI
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1639 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: N
ANTI-SENSE: N
FEATURE:
NAME/KEY: CDS
LOCATION: 126..1523
OTHER INFORMATION:
US-08-334-698-5

Query Match 2.7%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76; Mismatches 0; Gaps 0;
Matches 78; Conservative 0; Indels 72; Indels 0; Gaps 0;
QY 474 cgcgtccacggcctccagccacgggacacggcgtggtgcgttatctgcagcgctct 533
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTCTTCTCAGCGCT 694
QY 534 ggcacccgtctcgtggtgcgtctcctctgtctatctattgtaagagacagtttat 593
DB 695 GGGCTCTTCTACCTGCTCTGCGCATCATCTGCTGTCATGTACTGCGCGTCTACGTGCT 754
QY 594 ggaagaacacccagctgctctcgcgtc 623
DB 755 GCCAAGAGGGAGCGCGGGCTCAAGTC 784

RESULT 12
US-08-932-5
Sequence 5, Application US/08228932
Patent No. 5578611
GENERAL INFORMATION:
APPLICANT: Charles Gluchowski, Carlos C. Forray, George Chiu,
APPLICANT: Theresa A. Branche, John M. Wetzel and Paul R. Hartig
TITLE OF INVENTION: USE OF ALPHA-1C SPECIFIC COMPOUNDS TO TREAT BENIGN
TITLE OF INVENTION: PROSTATIC HYPERPLASIA
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM
STREET: 30 Rockefeller Plaza
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10112
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/228,932
FILING DATE: 13-APR-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: White, John P. 28,678
REGISTRATION NUMBER: 41878-B/JPW/TEP
REFERENCE/DOCKET NUMBER: 41878-B/JPW/TEP

TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEFAX: (212) 664-0525
TELEX: (212) 422523 COOP UI
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1639 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: N
ANTI-SENSE: N
FEATURE:
NAME/KEY: CDS
LOCATION: 126..1523
OTHER INFORMATION:
US-08-228-932-5

Query Match 2.7%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76; Mismatches 0; Gaps 0;
Matches 78; Conservative 0; Indels 72; Indels 0; Gaps 0;
QY 474 cgcgtccacggcctccagccacgggacacggcgtggtgcgttatctgcagcgctct 533
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTCTTCTCAGCGCT 694
QY 534 ggcacccgtctcgtggtgcgtctcctctgtctatctattgtaagagacagtttat 593
DB 695 GGGCTCTTCTACCTGCTCTGCGCATCATCTGCTGTCATGTACTGCGCGTCTACGTGCT 754
QY 594 ggaagaacacccagctgctctcgcgtc 623
DB 755 GCCAAGAGGGAGCGCGGGCTCAAGTC 784

RESULT 13
US-08-468-939-5
Sequence 5, Application US/08468939
Patent No. 5714381
GENERAL INFORMATION:
APPLICANT: Jonathan A. Bard et al.
TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
TITLE OF INVENTION: Receptors and Uses Thereof
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/468,939
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P. 28,678
REGISTRATION NUMBER: 41337-1B
REFERENCE/DOCKET NUMBER: 41337-1B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEFAX: (212) 391-0526
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1639 base pairs
TYPE: nucleic acid

		Matches	78;	Conservative	0;	Mismatches	72;	Indels	0;	Gaps	0;
Qy	474	cgcgtccacg	g	c	g	c	c	c	c	c	c
Db	635	CGAGGACG	G	A	C	C	G	A	C	C	G
Qy	534	ggccaccgt	c	c	t	c	t	c	t	c	t
Db	695	GGGCTCCT	T	T	A	C	T	G	C	T	T
Qy	594	ggagaagaa	a	a	a	a	a	a	a	a	a
Db	755	GGCCAAGG	A	G	G	A	G	A	G	A	G

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OM nucleic - nucleic search, using sw model

Run on: April 11, 2002, 00:23:18 ; Search time 107.54 seconds
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Title: US-09-380-276A-7
Perfect score: 1496
Sequence: 1 gggacgtagaactctccaa.....aaaaaaaaaaaaaaaaaaaaa 1496

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
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3: /cgn2_6/ptodata/2/ina/6A_COMB.seq.*
4: /cgn2_6/ptodata/2/ina/6B_COMB.seq.*
5: /cgn2_6/ptodata/2/ina/PTUS_COMB.seq.*
6: /cgn2_6/ptodata/2/ina/backfiles1.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	447.2	29.9	893	4	US-09-286-529-8
2	326.6	21.8	623	4	US-09-286-529-9
3	54.8	3.7	1582	3	US-08-545-196B-10
4	54.8	3.7	1582	3	US-08-545-196B-12
c 5	54.4	3.6	3138	1	US-07-867-106-4
6	53	3.5	2007	3	US-08-747-221B-36
c 7	53	3.5	2007	3	US-08-747-221B-38
8	53	3.5	2007	4	US-09-005-051-36
c 9	53	3.5	2007	4	US-09-005-051-38
10	52.8	3.5	5173	1	US-08-242-677-1
11	52.6	3.5	1420	2	US-08-909-965C-3
12	52.6	3.5	3238	4	US-08-123-934A-5
13	52.6	3.5	3238	5	PCT-US94-10080-5
14	52.4	3.5	3581	2	US-08-738-349-1
15	52	3.5	3709	4	US-09-541-782-7
16	51.8	3.5	3437	3	US-08-860-339-17
17	51.6	3.4	1641	1	US-08-300-903A-8
18	51.2	3.4	991	3	US-08-924-747-25
19	51.2	3.4	991	4	US-09-247-373B-25
20	51.2	3.4	991	4	US-09-296-715-25
21	51.2	3.4	1776	3	US-08-655-352-10
22	50.6	3.4	15124	2	US-08-487-826B-13
23	50.2	3.4	3933	1	US-08-199-776-1
24	50.2	3.4	3933	3	US-08-663-731-1
25	50.2	3.4	3933	3	US-08-879-338-1
26	50.2	3.4	3933	5	PCT-US95-02044-1
27	50	3.3	1046	1	US-08-361-467B-4

28	50	3.3	1046	1	US-08-484-332C-4	Sequence 4, Appli
29	50	3.3	1472	4	US-08-781-420-10	Sequence 10, Appl
c 30	50	3.3	1472	4	US-08-781-420-12	Sequence 12, Appl
31	49.8	3.3	8920	2	US-08-446-855A-1	Sequence 1, Appli
32	49.8	3.3	8920	4	US-09-150-741-1	Sequence 1, Appli
33	49.6	3.3	2700	3	US-09-315-861-1	Sequence 1, Appli
34	49.4	3.3	746	4	US-09-013-810-1	Sequence 1, Appli
35	49.2	3.3	2887	5	PCT-US96-10521-14	Sequence 14, Appl
36	49	3.3	5952	1	US-07-867-106-2	Sequence 2, Appli
37	48.8	3.3	458	1	US-08-524-757-1	Sequence 1, Appli
38	48.8	3.3	2469	4	US-09-111-730-5	Sequence 5, Appli
c 39	48.6	3.2	1582	3	US-08-545-196B-10	Sequence 10, Appl
c 40	48.6	3.2	1582	3	US-08-545-196B-12	Sequence 12, Appl
41	48.6	3.2	2589	4	US-08-569-749-1	Sequence 1, Appli
42	48.6	3.2	2589	5	PCT-US96-112860-1	Sequence 1, Appli
43	48.4	3.2	1393	1	US-08-174-467-18	Sequence 18, Appl
44	48.4	3.2	1393	3	US-08-452-071-18	Sequence 18, Appl
45	48.4	3.2	1490	2	US-08-553-367A-5	Sequence 5, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match	29.9%;	Score 447.2;	DB 4;	Length 893;
Best Local Similarity	81.6%;	Pred. No. 1.5e-101;		
Matches 529;	Conservative	0;	Mismatches 118;	Indels 1;
Gaps	1;			
QY	22	aataaatacatttgataaagaagatggctttaaaagtctactagacaagaagaaacgt	81	
Db	32	aataaacaagtttggtagagccatggcactaaagctcctctacacagagcaggtgc	91	
QY	82	ttttcactcttttagttactaggtctattgtcatgtaaaagtactgtgaaacaggag	141	
Db	92	tcttgcgtccattctctcctactccactcgcagtgatgagtgagtggaacccggag	151	
QY	142	actgtagacagaagaattcaggatcggtctgtggaactgtgttccctgcacacagtg	201	
Db	152	attgcaggcagcagggaattcaagatcgatctgtggaactgtgtcctctgcaaacag	211	
QY	202	gccacggcagtgagttgtctaaggaatgtggtctgcgctatgggaggaatgcacagt	261	
Db	212	gacctggcagtgagttgtccaaggaatgtggtctgcgctatgggaggaatgcacagt	271	
QY	262	tgacgtgcgcgctgcacaggttcaaggagactggggctccagaatgcagacccctgc	321	
Db	272	tgccctgcaggccgcaccsggttcaaggagactggggtttccagaagtgttaagccat	331	
QY	322	tggaactgcagtggtggaaccgctttcagaaggaataatgttccagccacagtgatgc	381	
Db	332	cggactgtgcgtggtagaccgctttcagaggcccaactgtcacacacagtgatgc	391	
QY	382	tctgcgggagactgctgcaggattttataggaagacgaactgtcgcgtttcaagaca	441	
Db	392	tctgcgggagactgctgcaggattttataggaagacgaactgtcgcgtttcaagaca	451	


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; Patent No. 6291222
; GENERAL INFORMATION:
; APPLICANT: Silver, Gary W.
; APPLICANT: Wisniewski, Nancy
; TITLE OF INVENTION: No. 6291222el Carboxylesterase Nucleic Acid
; TITLE OF INVENTION: Molecules, Proteins and Uses Thereof
; NUMBER OF SEQUENCES: 66
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carol Talkington Verser, Ph.D.
; ADDRESSEE: Heska Corporation
; STREET: 1825 Sharp Point Drive
; CITY: Fort Collins
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80525
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 95
; SOFTWARE: WordPerfect for Windows, Version 7.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/005,051
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/747,221
; FILING DATE: No. 6291222ember 12, 1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Verser, Carol Talkington
; REGISTRATION NUMBER: 37,459
; REFERENCE/DOCKET NUMBER: FC-1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 970/493-7272
; TELEFAX: 970/484-9505
; INFORMATION FOR SEQ ID NO: 38:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2007 nucleotides
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; US-09-005-051-38

Query Match 3.5%; Score 53; DB 4; Length 2007;
Best Local Similarity 67.9%; Pred. No. 0.00064;
Matches 74; Conservative 0; Mismatches 35; Indels 0; Gaps 0;

Qy 1384 ttttttttgcattcttaataattcttgtagtgtagagtagtcttttaaaataaatt 1443
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 109 ttattaccatctttgtatcatatatttgcatttttttttttttttttttttttttcaata 50
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Qy 1444 tcaagtatttttttaaaactaaataaaataaaataaaataaaataaaataaaataaa 1492
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 49 ttttttttttttttttttttttttttttttttttttttttttttttttttttttttttt 1
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |

RESULT 10
US-08-242-677-1
; Sequence 1, Application US/08242677
; Patent No. 5677143
; GENERAL INFORMATION:
; APPLICANT: Gaynor, Richard B
; APPLICANT: Wu, Foon W.
; TITLE OF INVENTION: Cellular Nucleic Acid Binding Protein
; TITLE OF INVENTION: and Uses Thereof in regulating Gene Expression and in the
; TITLE OF INVENTION: Treatment of AIDS
; NUMBER OF SEQUENCES: 9
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Arnold, White & Durkee
; STREET: P.O. Box 4433
; CITY: Houston
; STATE: TX
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; COUNTRY: USA
; ZIP: 77210
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/242,677
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Mayfield, Denise L.
; REGISTRATION NUMBER: 33,732
; REFERENCE/DOCKET NUMBER: UTSD:401
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 713-787-1400
; TELEFAX: 713-789-2679
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5173 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 1..4863
; US-08-242-677-1

Query Match 3.5%; Score 52.8; DB 1; Length 5173;
Best Local Similarity 67.0%; Pred. No. 0.0011;
Matches 75; Conservative 0; Mismatches 37; Indels 0; Gaps 0;

Qy 1385 ttttttttgcattctttaaataattcttgtagtgtagagtagtcttttaaaataaatt 1444
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 5028 TATTTTCCCTTTAATAAACACTTTTGTAAATGTATCTCTTCTTAAATAAATATT 5087
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Qy 1445 caagtatttttttaaaactaaataaaataaaataaaataaaataaaataaaataaa 1496
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 5088 TAAGCAATTGTCCATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAA 5139
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |

RESULT 11
US-08-909-965C-3
; Sequence 3, Application US/08909965C
; Patent No. 5936078
; GENERAL INFORMATION:
; APPLICANT: Kuga Tetsuo
; APPLICANT: Nakagawa Satoshi
; APPLICANT: Sakaki Yoshiyuki
; APPLICANT: Zhao Nanding
; APPLICANT: Hashida Hideji
; TITLE OF INVENTION: NOVEL DNA, NOVEL POLYPEPTIDE
; TITLE OF INVENTION: AND NOVEL ANTIBODY
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FITZPATRICK, CELLA, HARPER AND SCINTO
; STREET: 277 Park Avenue
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10172-0194
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/909,965C
; FILING DATE: August 12, 1997
; CLASSIFICATION: 514
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; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
; IMMEDIATE SOURCE:
; CLONE: CFK1-10a
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 474..2000
PCT-US94-10080-5

Query Match      3.5%; Score 52.6; DB 5; Length 3238;
Best Local Similarity 68.2%; Pred. No. 0.001;
Matches 73; Conservative 0; Mismatches 34; Indels 0; Gaps 0;

QY 1390 ttttgcattttaaattcttgcattgtgtgtagagtagtgtttttaaataatttcaagt 1449
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Db 3091 TATTTGTTTAACTACTTTTGTATTAGTAGTATTATTGTATAAATTAATAAACTGT 3150
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 1450 atttttttaaaactaaataaaataaaataaaataaaataaaataaaataaaataaa 1496
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 3151 TTCAAGTCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 3197

RESULT 14
US-08-738-349-1
; Sequence 1, Application US/08738349
; Patent No. 5869638
; GENERAL INFORMATION:
; APPLICANT: Takeshita, Sunao
; APPLICANT: Okazaki, Makoto
; APPLICANT: Kawai, Shinji
; APPLICANT: Tsujimura, Atsushi
; APPLICANT: Amann, Egon
; TITLE OF INVENTION: Bone-Related Cadherin-Like Protein and
; TITLE OF INVENTION: Process for Its Production
; NUMBER OF SEQUENCES: 12
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; ADDRESSEE: Dunner
; STREET: 1300 I Street, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005-3315
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/738,349
; FILING DATE: 25-OCT-1996
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/364,439
; FILING DATE:
; APPLICATION NUMBER: US 08/112,061
; FILING DATE: 26-AUG-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Barker, M. P.
; REGISTRATION NUMBER: 32,013
; REFERENCE/DOCKET NUMBER: 02481.1323-00000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-408-4000
; TELEFAX: 202-408-4400
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3581 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA to mRNA
; ORIGINAL SOURCE:

; ORGANISM: Mus musculus
; STRAIN: osteoblastic cell line MC3T3E1
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 284..2671
US-08-738-349-1

Query Match      3.5%; Score 52.4; DB 2; Length 3581;
Best Local Similarity 67.3%; Pred. No. 0.0012;
Matches 74; Conservative 0; Mismatches 36; Indels 0; Gaps 0;

QY 1387 ttttttgcattttaaattcttgcattgtgtgtagagtagtgtttttaaataatttca 1446
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Db 3434 TTGTACTGTGTCTTTTAATATGAGCTTCAATATAAGAAGCAAGCTTTTGAATAAAAAAAG 3493
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 1447 agtattttttttaaactaaataaaataaaataaaataaaataaaataaaataaa 1496
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 3494 ATTCCTTTTAAAAAATAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 3543

RESULT 15
US-09-541-782-7
; Sequence 7, Application US/09541782
; Patent No. 6284480
; GENERAL INFORMATION:
; APPLICANT: Nislow, Corey
; APPLICANT: Sakowicz, Roman
; APPLICANT: Beraud, Christophe
; TITLE OF INVENTION: Antifungal Assay
; FILE REFERENCE: 1015
; CURRENT APPLICATION NUMBER: US/09/541,782
; CURRENT FILING DATE: 2000-04-03
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 7
; LENGTH: 3709
; TYPE: DNA
; ORGANISM: Drosophila melanogaster
US-09-541-782-7

Query Match      3.5%; Score 52; DB 4; Length 3709;
Best Local Similarity 62.1%; Pred. No. 0.0015;
Matches 82; Conservative 0; Mismatches 50; Indels 0; Gaps 0;

QY 1365 ttgggggaacctgatgagtttttttttttgcatctttaaataattcttctgtgtgtaga 1424
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 3572 ttgtttggctctcttggtatctcatcttctgtatcatctcttggaattgtgaca 3631
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 1425 gtatgtttttaaataaaatttcaagtattttttttaaactaaataaaataaaataaa 1484
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 3632 ttattattttaaataaaactcagtagtctatgtaaaagttaaataaaataaaataaa 3691
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 1485 aaaaaaaaaaaaaa 1496
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 3692 aaaaaaaaaaaaaa 3703
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Search completed: April 11, 2002, 00:23:32
Job time: 10279 sec
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GenCore version 4.5
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OM protein - protein search, using sw model

Run on: April 10, 2002, 17:16:24 ; Search time 21.63 Seconds
(without alignments)
440.078 Million cell updates/sec

Title: US-09-380-276a-8
Perfect score: 2283
Sequence: 1 MALKVLEQEKFTFTLLVLL.....AIIHPATQTSIQVRQRLGSL 423

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 212252 seqs, 22503292 residues

Total number of hits satisfying chosen parameters: 212252

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep.*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep.*
5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep.*
6: /cgn2_6/ptodata/2/iaa/backfiles1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	869	38.1	210	4	US-09-286-529-3
2	714.5	31.3	151	4	US-09-286-529-4
3	159	7.0	438	1	US-08-097-827-11
4	159	7.0	438	1	US-08-494-574-11
5	150	6.6	206	1	US-08-097-827-7
6	150	6.6	206	1	US-08-494-574-7
7	145	6.4	205	3	US-08-974-022-51
8	145	6.4	205	4	US-08-795-445A-51
9	145	6.4	205	4	US-08-795-447A-51
10	145	6.4	205	4	US-08-974-186-51
11	145	6.4	205	4	US-08-795-446B-51
12	144	6.3	1104	2	US-08-327-832-5
13	144	6.3	1104	2	US-08-828-584-5
14	136	6.0	625	3	US-08-996-139-15
15	136	6.0	625	4	US-08-995-659-15
16	136	6.0	625	4	US-09-215-649A-15
17	134.5	5.9	415	4	US-09-006-353A-6
18	134	5.9	186	1	US-08-089-458B-6
19	134	5.9	307	4	US-08-804-166-4
20	134	5.9	307	4	US-08-910-991-4
21	133.5	5.8	2050	2	US-08-347-594A-2
22	132.5	5.8	197	2	US-08-505-606-1
23	132	5.8	276	4	US-09-041-886-27
24	132	5.8	277	4	US-09-042-785A-10
25	132	5.8	277	4	US-09-006-353A-10
26	131	5.7	139	2	US-08-219-237B-8
27	131	5.7	176	4	US-09-411-722-1

28	130.5	5.7	140	4	US-08-477-347-17	Sequence 17, Appl
29	130.5	5.7	140	4	US-08-476-862-8	Sequence 8, Appli
30	129.5	5.7	336	4	US-08-804-166-8	Sequence 8, Appli
31	129.5	5.7	336	4	US-08-910-991-8	Sequence 8, Appli
32	129	5.7	326	1	US-08-292-549-4	Sequence 4, Appli
33	129	5.7	326	5	PCT-US91-02207-4	Sequence 4, Appli
34	128.5	5.6	1170	1	US-08-313-288B-20	Sequence 20, Appl
35	128	5.6	355	1	US-08-292-549-6	Sequence 6, Appli
36	128	5.6	355	4	US-09-006-353A-14	Sequence 14, Appl
37	127	5.6	419	4	US-08-509-024-7	Sequence 7, Appli
38	127	5.6	419	4	US-09-333-279-7	Sequence 7, Appli
39	126	5.5	148	4	US-09-411-722-2	Sequence 2, Appli
40	124.5	5.5	1111	1	US-08-317-450B-15	Sequence 15, Appl
41	124.5	5.5	1111	4	US-08-800-593-15	Sequence 15, Appl
42	124.5	5.5	1193	1	US-08-317-450B-13	Sequence 13, Appl
43	124.5	5.5	1193	4	US-08-800-593-13	Sequence 13, Appl
44	124.5	5.5	2813	3	US-08-896-449A-2	Sequence 2, Appli
45	124.5	5.5	2813	3	US-09-132-652-2	Sequence 2, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-3
; Sequence 3, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catharine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 210
; TYPE: PRT
; ORGANISM: human
US-09-286-529-3

Query Match 38.1%; Score 869; DB 4; Length 210;
Best Local Similarity 83.7%; Pred. No. 3.le-73;
Matches 154; Conservative 11; Mismatches 19; Indels 0; Gaps 0;

QY	1	MALKVLEQEKFTFTLLVLLGYLSCKVTCETGDCRQOEPRDRSGNCVPCNCGPMELSK	60
Db	1	MALKVLP LHTVLF AAILFLHLACKVSCETGDCRQOEFKDRSGNCVLCKQCGPMELSK	60
QY	61	ECGFGYGEDAQCVTCRLHREFKEDWGFKCKPCLDCAVVRFRKANGSATSDAICGDCPLG	120
Db	61	ECGFGYGEDAQCVCPHRFEKEDWGFKCKPCADCALVRFRANCSTSDAVCGDCPLG	120
QY	121	FYRKTKLVGFQDMCEVPCGDP PPPPPYPHPCASKVNLVKIATASPRDTAAAVICSALAT	180
Db	121	FYRKTKLVGFQDMCEVPCGDP PPPPPYPHCTSKVNLVKISSTVSSPRDTAAAVICSALAT	180
QY	181	VLLA 184	
Db	181	VLLA 184	

RESULT 2
US-09-286-529-4
; Sequence 4, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catharine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05

NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 151
; TYPE: PRT
; ORGANISM: human
US-09-286-529-4

Query Match 31.3%; Score 714.5; DB 4; Length 151;
Best Local Similarity 82.0%; Pred. No. 4.6e-59;
Matches 123; Conservative 9; Mismatches 17; Indels 1; Gaps 1;

QY 1 MALKVLEQEKTFETLLVLLGYLSCKVTCTGDC-RQQRDRSGNCPVPCNQCQGMELSG 59
DB 1 MALKVLPRLHRTVFAAILLLHACKVSCETGDCSRQQEFKDRSGNVLCKQCGPMELSG 60
QY 60 KECFGYGEDAQCVCTCLHREFKEDWGQKPCLDCAVNRFOKANCATSATSDAICGDCLP 119
DB 61 KECFGYGEDAQCVCPKPHREFKEDWGQKPCADCALVNRFORANCSTSDAICGDCLP 120
QY 120 GFYRKTKLVGFQDMCVPCGDPPTPPPEPHC 149
DB 121 GFYRKTKLVGFQDMCVPCGDPPTPPPEPHC 150

RESULT 3

US-08-097-827-11

; Sequence 11, Application US/08097827

; GENERAL INFORMATION:

; APPLICANT: Baum, Peter

; Goodwin, Ray

; Fanslow, William

; Gayle, Richard

; TITLE OF INVENTION: Novel Cytokine which is a Ligand for

; OX40

; NUMBER OF SEQUENCES: 13

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Immunex Corporation

; STREET: 51 University Street

; CITY: Seattle

; STATE: WA

; COUNTRY: USA

; ZIP: 98101

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/097,827

; FILING DATE: 23-Jul-1993

; CLASSIFICATION: <Unknown>

; ATTORNEY/AGENT INFORMATION:

; NAME: Perkins, Patricia A.

; REGISTRATION NUMBER: 34,693

; REFERENCE/DOCKET NUMBER: 2806

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 206-587-0730

; INFORMATION FOR SEQ ID NO: 11:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 438 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; SEQUENCE DESCRIPTION: SEQ ID NO: 11:

US-08-097-827-11

Query Match

Best Local Similarity 7.0%; Score 159; DB 1; Length 438;

Matches 69; Conservative 31; Mismatches 93; Indels 58; Gaps 18;

QY 15 TLLVLLGYLSCKVTCTGDCRQQRDRSGN-CVPCNQCQGMELSGKECGFYGEDAQCV 73
DB 9 TALLLLG-LTLGVTARRLNCVKHTY--PSGKHC--CRECOPGHGMVNR--DHTRTLCH 61
QY 74 TCRHREFKEDWGQKPCLDCAVNRFO-KANCATSATSDAICGDCLPGFYRKTKLVGFQD 132
DB 62 PCETGFYNEAVNYDTCKQCTQCNRHSGSELKQCTPTQDTVC-RCRPGTQPR-----QD 114
QY 133 -----MECVPCGDPPTPPPEP-----HCASKVNLVKLTASTASSPRDTALAIVC---SALA 179
DB 115 SGYKLVGDCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSLDV-CEDRSLLA 171
QY 180 TVLLALLLVCIVYCKRQFMEKKPSW---SLRSQDIQVNGSELSCLDPRPQLHEVAHRACQ 236
DB 172 TLL-----WETQRTFRTTVQSTTVMPRTSELP--STPTLVE--PRSC-- 211
QY 237 CRDSVOTCGP 247
DB 212 ---DKTHTCPT 219

RESULT 4

US-08-494-574-11

; Sequence 11, Application US/08494574

; Patent No. 5783665

; GENERAL INFORMATION:

; APPLICANT: Baum, Peter

; APPLICANT: Goodwin, Ray

; APPLICANT: Fanslow, William

; APPLICANT: Gayle, Richard

; TITLE OF INVENTION: No. 5783665el Cytokine which is a Ligand for

; NUMBER OF SEQUENCES: 13

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Immunex Corporation

; STREET: 51 University Street

; CITY: Seattle

; STATE: WA

; COUNTRY: USA

; ZIP: 98101

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/494,574

; FILING DATE: 22-JUN-1995

; CLASSIFICATION: 530

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US/08/097,827

; FILING DATE: 23-JUL-1993

; ATTORNEY/AGENT INFORMATION:

; NAME: Perkins, Patricia A.

; REGISTRATION NUMBER: 34,693

; REFERENCE/DOCKET NUMBER: 2806

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 206-587-0730

; INFORMATION FOR SEQ ID NO: 11:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 438 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-494-574-11

Query Match

Best Local Similarity 7.0%; Score 159; DB 1; Length 438;

Matches 69; Conservative 31; Mismatches 93; Indels 58; Gaps 18;

QY 15

TLLVLLGYLSCKVTCTGDCRQQRDRSGN-CVPCNQCQGMELSGKECGFYGEDAQCV 73

Db 9 TALLLLG-LTLGVTTARRLNCVKHTY--PSGHKC--CRECQPGHGMVNR--DHTRDTLCH 61

QY 74 TCLRHRFKEDMGFOKCKPLCDCAVNVNRQ-KANCATSDAICGDCPLPGFYRKTKLVGFQD 132

Db 62 PCETGFYNEAVNYDTCKQCTQCNHRSGSELKQNCPTQDQTV--RCRPGTQPR-----QD 114

QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIASTASSPRDTALAAVIC-----SALA 179

Db 115 SGYKLGVDVCVC--PPGHFSPGNNOACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171

QY 180 TVL 182

Db 172 TLL 174

RESULT 6

US-08-494-574-7

; Sequence 7, Application US/08494574

; Patent No. 5783665

; GENERAL INFORMATION:

; APPLICANT: Baum, Peter

; APPLICANT: Goodwin, Ray

; APPLICANT: Fanslow, William

; APPLICANT: Gayle, Richard

; TITLE OF INVENTION: No. 5783665el Cytokine Which is a Ligand for

; TITLE OF INVENTION: OX40

; NUMBER OF SEQUENCES: 13

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Immunex Corporation

; STREET: 51 University Street

; CITY: Seattle

; STATE: WA

; COUNTRY: USA

; ZIP: 98101

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/494,574

; FILING DATE: 22-JUN-1995

; CLASSIFICATION: 530

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US/08/097,827

; FILING DATE: 23-JUL-1993

; ATTORNEY/AGENT INFORMATION:

; NAME: Perkins, Patricia A.

; REGISTRATION NUMBER: 34,693

; REFERENCE/DOCKET NUMBER: 2806

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 206-587-0730

; INFORMATION FOR SEQ ID NO: 7:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 206 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; SEQUENCE DESCRIPTION: SEQ ID NO: 7:

US-08-097-827-7

Query Match 6.6%; Score 150; DB 1; Length 206;

Best Local Similarity 30.1%; Pred. No. 2.4e-06;

Matches 55; Conservative 21; Mismatches 75; Indels 32; Gaps 13;

QY 15 TLLVLLGLYSCKVTCTGDCRQOEFRDRSGN-CVPCNQCGPMGSMELSKGCGFYGEDAQCV 73

Db 9 TALLLLG-LTLGVTTARRLNCVKHTY--PSGHKC--CRECQPGHGMVNR--DHTRDTLCH 61

QY 74 TCLRHRFKEDMGFOKCKPLCDCAVNVNRQ-KANCATSDAICGDCPLPGFYRKTKLVGFQD 132

Db 62 PCETGFYNEAVNYDTCKQCTQCNHRSGSELKQNCPTQDQTV--RCRPGTQPR-----QD 114

QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIASTASSPRDTALAAVIC-----SALA 179

Db 115 SGYKLGVDVCVC--PPGHFSPGNNOACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171

QY 180 TVL 182

QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIASTASSPRDTALAAVIC-----SALA 179

Db 115 SGYKLGVDVCVC--PPGHFSPGNNOACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171

QY 180 TVL 182

Db 172 TLL 174

RESULT 6

US-08-494-574-7

; Sequence 7, Application US/08494574

; Patent No. 5783665

; GENERAL INFORMATION:

; APPLICANT: Baum, Peter

; APPLICANT: Goodwin, Ray

; APPLICANT: Fanslow, William

; APPLICANT: Gayle, Richard

; TITLE OF INVENTION: No. 5783665el Cytokine Which is a Ligand for

; TITLE OF INVENTION: OX40

; NUMBER OF SEQUENCES: 13

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Immunex Corporation

; STREET: 51 University Street

; CITY: Seattle

; STATE: WA

; COUNTRY: USA

; ZIP: 98101

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/494,574

; FILING DATE: 22-JUN-1995

; CLASSIFICATION: 530

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US/08/097,827

; FILING DATE: 23-JUL-1993

; ATTORNEY/AGENT INFORMATION:

; NAME: Perkins, Patricia A.

; REGISTRATION NUMBER: 34,693

; REFERENCE/DOCKET NUMBER: 2806

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 206-587-0730

; INFORMATION FOR SEQ ID NO: 7:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 206 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; US-08-494-574-7

Query Match 6.6%; Score 150; DB 1; Length 206;

Best Local Similarity 30.1%; Pred. No. 2.4e-06;

Matches 55; Conservative 21; Mismatches 75; Indels 32; Gaps 13;

QY 15 TLLVLLGLYSCKVTCTGDCRQOEFRDRSGN-CVPCNQCGPMGSMELSKGCGFYGEDAQCV 73

Db 9 TALLLLG-LTLGVTTARRLNCVKHTY--PSGHKC--CRECQPGHGMVNR--DHTRDTLCH 61

QY 74 TCLRHRFKEDMGFOKCKPLCDCAVNVNRQ-KANCATSDAICGDCPLPGFYRKTKLVGFQD 132

Db 62 PCETGFYNEAVNYDTCKQCTQCNHRSGSELKQNCPTQDQTV--RCRPGTQPR-----QD 114

QY 133 -----MECVPCGDDPPPPYEP-----HCASKVNLVKIASTASSPRDTALAAVIC-----SALA 179

Db 115 SGYKLGVDVCVC--PPGHFSPGNNOACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171

QY 180 TVL 182

Db 172 TLL 174

RESULT 7

US-08-974-022-51
; Sequence 51, Application US/08974022
; Patent No. 6015938
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Behavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/974,022
; FILING DATE: 12-DEC-1995
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-974-022-51

Query Match 6.4%; Score 145; DB 3; Length 205;

Best Local Similarity 28.6%; Pred. No. 6.9e-06;

Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QEKTFLLVLLGLYLSCKVTCGTGDCRQOEFRDR---SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QOPTAFLLGLSLGVTVKLNC-----VKDTYPSGHKC--CRECPGHGMVSRD--D 52
QY 66 YGEDAQCVCRLHREKEDMGFKCKPCLDCAVVNRFO-KANCSDAICGDCPLCPGYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKOCTCNHRSGSELKQNCPTEDTVC-QCRPGTQPR 111
QY 125 TKLVGFQDMCEVPCGPPPPYEP-----HCASKYNLV-----KIASTASSPRDTALAAVIC- 175
Db 112 QDSSHLKGVDCVPC---PPGHFSPGSGNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164
QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 8

US-08-795-445A-51
; Sequence 51, Application US/08795445A
; Patent No. 6284485
; GENERAL INFORMATION:

; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Behavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,445A
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-795-445A-51

Query Match 6.4%; Score 145; DB 4; Length 205;

Best Local Similarity 28.6%; Pred. No. 6.9e-06;

Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QEKTFLLVLLGLYLSCKVTCGTGDCRQOEFRDR---SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QOPTAFLLGLSLGVTVKLNC-----VKDTYPSGHKC--CRECPGHGMVSRD--D 52
QY 66 YGEDAQCVCRLHREKEDMGFKCKPCLDCAVVNRFO-KANCSDAICGDCPLCPGYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKOCTCNHRSGSELKQNCPTEDTVC-QCRPGTQPR 111
QY 125 TKLVGFQDMCEVPCGPPPPYEP-----HCASKYNLV-----KIASTASSPRDTALAAVIC- 175
Db 112 QDSSHLKGVDCVPC---PPGHFSPGSGNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164
QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 9

US-08-795-447A-51
; Sequence 51, Application US/08795447A
; Patent No. 6284728
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: Osteoprotegerin
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: One Amgen Center Drive

```

: CITY: Thousand Oaks
: STATE: California
: COUNTRY: USA
: ZIP: 91362-1789
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: Patent In Release #1.0, Version #1.30
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/08/795,447A
: FILING DATE:
: CLASSIFICATION: 514
: ATTORNEY/AGENT INFORMATION:
: NAME: Winter, Robert B.
: REFERENCE/DOCKET NUMBER: A-378D2
: INFORMATION FOR SEQ ID NO: 51:
: SEQUENCE CHARACTERISTICS:
: LENGTH: 205 amino acids
: TYPE: amino acid
: STRANDEDNESS: single
: TOPOLOGY: linear
: MOLECULE TYPE: protein
: US-08-795-447A-51

```

```

Query Match      6.4%; Score 145; DB 4; Length 205;
Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QEKFTFTLLVLLGYSCKVTCTGDCRQEQPRDR--SGN-CVPCNQCQPGMGLSKECGFG 65
   | | | | | : | : | : | | : | | : | | : | | : |
Db 6 QQPTAFLLGLSLGVTYKLN-----VKDIYPSGHC--CRECQPGHGMVSR--D 52

QY 66 YGEDAQCVTCRLHREFKEDWGQKPCPLDCAVARNFQ-KANCATSDAICGDCLPGFYRK 124
   : | : | : | : | : | : | : | : | : | : | : | :
Db 53 HTRDTVCHPCPGFYNEAVNYDTCKQCTQCNRHSGSELKQNCPTPTDTCV--QCRPGTQPR 111

QY 125 TKLVGFQDMCVCGDPPPPYEP----HCASKVNLV----KIATASSPROTALAAVTC- 175
   : : : : : | | | : | : | : | : | : | : | : |
Db 112 QDSHKLGVDCVPC--PPGHFSPGSNACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164

QY 176 --SALATVL 182

Db 165 DRSLATLL 173

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```

RESULT 10
US-08-974-186-51
; Sequence 51, Application US/08974186
; Patent No. 628470
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/974,186
; FILING DATE:

```

```

; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-974-186-51

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Query Match	6.4%;	Score 145;	DB 4;	Length 205;
Best Local Similarity	28.6%;	Pred. No. 6.9e-06;		
Matches	54;	Conservative 23;	Mismatches 76;	Indels 36; Gaps 12;
Qy	9	QEKFTFTLLVLGLSCKVTCTGDCRQQRFRDR--SGN-CVPCNQCSPGMELSKCEGFG	65	
Db	6	QQPTAFLLGLSLGLGVTKLNC-----VKDIYPSGHCK--CRECQPGHGMVSRCC--D	52	
Qy	66	YGEDAQCVCYTKLHRFKEDWGFQKPCPLDCAVNRFO-KANCSTASDAICGDCLPGFYRK	124	
Db	53	HTRDTVCHPCPFGFYNEAVNYDTCQCTQCNRHSGSELKQNCPTEDTIVC--QCRPGTQPR	111	
Qy	125	TKLVGFQDMECVPCGDPDPYYP--HCASKVNLV---KIASTSPRDTALAAVTC-	175	
Db	112	QDSHKLGVDCVPC--PPGHFSPGNCQCKPWTNCTLSGKQIRHPASNSLDT-----VCE	164	
Qy	176	--SALATVL	182	
Db	165	DRSLLATLL	173	

```

RESULT" 11
US-08-795-446B-51
; Sequence 51, Application US/08795446B
; Patent No. 6288032
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patencin Release
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,446B
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:

```

```

; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-795-446B-51

```

```

Query Match      6.4%; Score 145; DB 4; Length 205;
Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QEKFTFTLLVLGLYSCKVTCTGDCRQEQPRDR--SGN-CVPNCQCGPMGSLKECGFG 65
   | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 6 QOPTAFLLGLSLGVTVKLC-----VKDTYPSGHC--CRECPGHGWSRC--D 52

QY 66 YGEDAQCVTCRLHRFKEDMGFKCPCLDCAVNRFO-KANCSTATSDAICGDCLPFGFYRK 124
   | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 53 HTRDTVCHPCGPGFYNEAVNYDTCKQCTQHRSSELSLKQCTPTEDTVC--QCRPGTQPR 111
   | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
QY 125 TKLVGFQDMECVPCGDP PPPYEP-----HCASKVNLV---KIATSPRDTALAAVIC- 175
   | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 112 QDSHKLGLVDCVFC--PPGHFSPGSCNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164

QY 176 --SALATVL 182
   | | | | |
Db 165 DRSLLATLL 173

```

```

RESULT 12
US-08-327-832-5
; Sequence 5, Application US/08327832
; Patent No. 5840832
; GENERAL INFORMATION:
; APPLICANT: Ono, Santa J.
; APPLICANT: Strominger, Jack L.
; TITLE OF INVENTION: Transcription Factor Regulating MHC
; TITLE OF INVENTION: Expression, cDNA and Genomic Clones Encoding Same and
; TITLE OF INVENTION: Retroviral Expression Constructs Thereof
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Banner, Birch, McKie & Beckett
; STREET: 1001 G Street, N.W.
; CITY: Washington, D.C.
; STATE: District of Columbia
; COUNTRY: U.S.A.
; ZIP: 20001
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/327,832
; FILING DATE:
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: Posorske, Laurence H.
; REGISTRATION NUMBER: 34,698
; REFERENCE/DOCKET NUMBER: 1107.46362
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 20-2 508-9153
; TELEFAX: 202 508-9299
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1104 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-327-832-5

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Query Match

	Best Local Similarity	18.8%;	Pred. NO.	9e-05;	
	Matches	110;	Conservative	59;	Mismatches 181; Indels 236; Gaps 27;
QY	24	SCKVTCTGDCRQOERDRSGNVPQCNCQCPGMELSKECG-----FGYGEDAQC----	72		
DB	440	SCNLCHPG-----PCPPCAFAETKTCECGTRHTVRCGQAIVHCSNPC	484		
QY	73	---VTCRLHREFKDWDGFKCKPCLDCAVNRFKANCATS--DAICGDCLPGPYRKTKLV	128		
DB	485	ENTLNCQHGOAEIHCGGOGCOPCO--IILN~-QVCYGGSTSRVLCTGDV-----GKSD	534		
QY	129	GFODMEG-----VPCGD-----PPP-----PYEPHC--ASKVNLVKIASTASS	164		
DB	535	GFGDFSLCKTGCGLDKCGNITCSQVCHPQCQCQPRLPQLVRCPCGQTPLSOLLELGSS	594		
QY	165	PRDTALAAV-----IC-SALATVLLALLILC-----VIYCKRFMEKK-	201		
DB	595	SRKTCMDPVPCGVCKGPLPCGSLDIHTCEKLCHGEDGCPVSRTSVISCRGSFRTKEL	654		
QY	202	PSWLSRSODI-----QYNGSELSCLDLRPO-----LH---	EYA	230	
DB	655	PTSLKSSEDATFCMCDKRCNKRRKLCGRHKCNIEICCVDEKHKCPLNCKRLRCGLHRCEEPC	714		
QY	231	HRACCQ-CRRDSVQT-----CGPVRLLPMSOC-----EE	258		
DB	715	HRGNQCICWAASFDLTCHCGASVIYPVPCGTTPPECTQCARVHECDHPVYHSGSEE	774		
QY	259	ACS-----PNPATIGCGVHSAASL-----	277		
DB	775	KCPPCTFLTQKWCMKGKHEFRSNIPCHLVDISCGLPUSATLPCGMHKKCORLCHKGECLVDE	834		
QY	278	-----QAR-----NAGPAGEMVPTFEGLTSQSI	300		
DB	835	PKQPCTTPRADCGHPCMARCHTSSPCPVTAACKAVELOCCEGRRKEWJCSBASSTYQR	894		
QY	301	CGEFSDAWLQNPMPGMDNISFCDSYPELTGEDITHSLNPELESSTLSDNSSQDLVGAV	360		
DB	895	IAAISMASKITDMLGGS-----VEISKILTKEVHQARLECDCECSALERKKR--LAEAF	948		
QY	361	PVOHSSENFATAATDLSRYNTLVESASTODALTMRSSOLDQESNAII	406		
DB	949	HIESDSPFMIRSGSKSFSDSLKEDA--RKDLKFVSDVERKEMETLV	992		
 RESULT 13 US-08-828-584-5 Sequence 5, Application US/08828584 Patent No. 5908762 GENERAL INFORMATION: APPLICANT: Ono, Santa J. TITLE OF INVENTION: Strominger, Jack L. TITLE OF INVENTION: Transcription Factor Regulating MHC TITLE OF INVENTION: Expression, cDNA and Genomic Clones Encoding Same and TITLE OF INVENTION: Retroviral Expression Contracts Thereof NUMBER OF SEQUENCES: 16 CORRESPONDENCE ADDRESS: ADDRESSEE: Banner, Birch, McKie & Beckett STREET: 1001 G Street, N.W. CITY: Washington, D.C. STATE: District of Columbia COUNTRY: U.S.A. ZIP: 20001 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/828,584 FILING DATE: CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION: NAME: Posorske, Laurence H.					

REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 1107.46362
TELECOMMUNICATION INFORMATION:
TELEPHONE: 20-2 508-9153
TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1104 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-828-584-5

Query Match 6.3%; Score 144; DB 2; Length 1104;
Best Local Similarity 18.8%; Pred. No. 9e-05;
Matches 110; Conservative 59; Mismatches 181; Indels 236; Gaps 27;

QY 24 SKVTCETGDCRQEQFRDRSGNVCNCGCPGMELSKGCG-----FGYGEDAOC----- 72
DB 440 SCNLLCHPG-----PCPPCPAFMTKCEGGRTRHTVRCQAVSVHCSNPC 484
QY 73 ---VTCRLHREKEDGFKCKPCDCAVNVRFQKANCATS-DAICGDCPLPGFYRKTLV 128
DB 485 ENILNCGQHCAELCHGGCQCPCQ--IILN--QVCYCGSTSRDVLGTDV-----GKSD 534
QY 129 GFQDMEC-----VPCGD-----PPP-----PYEPHC--ASKVNLVKIASTASS 164
DB 535 GDFSCILKTCGKDLKCGNHTSCVCHQPCQCPRLPOLVRCPCGQTPPLSLLLELGS 594
QY 165 PRDTALAAV-----IC-SALATVILALLILC-----VYICKRQFMKK- 201
DB 595 SRKTCMDPVPCGKVCCKPLCGSLDFIHTCEKLCHEGDCGPVSRVTSVISCRCSPRTKEL 654
QY 202 PSWLSRSQDI-----QYNGSELSCLDLDPQ-----LH-----EVA 230
DB 655 PCTLSKSEDATFMCDCRKNKRLCGRHKNEICCVDKHCKPLNCGRLKRLCGLHRCCEPC 714
QY 231 HRACCO-CRRDSVQT-----CGVRLLLPSMCC-----EE 258
DB 715 HRGNCQTCWQASFDLTCGASVIYPPVPCGTRPPECQTCTCARVHECDHPVYHSHSE 774
QY 259 ACS-----PNPATLGCYVHSAASL----- 277
DB 775 KCPPCTFLTKQCMGKHFEPSNIPCHLVDISGLPCSATLPCGMHKKCORLCHKGECLVDE 834
QY 278 -----QAR-----NAGPAGEMVPTFFGSLTQSI 300
DB 835 PCKQPCITPRADGHPCMAPHTSSPCPVYACKAKVLOCEGRRKEMVICSEASTYOR 894
QY 301 CGEFSDAWPLMNPMDGNISFCDSPYELTGEDIHSLNPELSSTSLDSNSQDLVGGAV 360
DB 895 IAAISMASKITDMLGGS-----VEISKLTIKKEVHQARLECECSALERKKR--LAEEF 948
QY 361 PVQSHENFTATDLSRYNNTLVESASTODALTMRSDQESGAI 406
DB 949 HISESDPENIRSSGKSFSDSLKEDA--RKDLKFVSDVEKMETLV 992

RESULT 14
US-08-996-139-15
Sequence 15, Application US/08996139
Patent No. 6017729
GENERAL INFORMATION:
APPLICANT: Anderson, Dirk M.
APPLICANT: Galibert, Laurent
APPLICANT: Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSER: Immunex Corporation, Law Department
STREET: 51 University Street
CITY: Seattle

STATE: WA
COUNTRY: USA
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/996,139
FILING DATE: 22 DECEMBER 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 60/064,671
FILING DATE: 14 OCTOBER 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2851-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 625 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-996-139-15

Query Match 6.0%; Score 136; DB 3; Length 625;
Best Local Similarity 22.2%; Pred. No. 0.00023;
Matches 102; Conservative 51; Mismatches 171; Indels 136; Gaps 27;

QY 16 LVLVLGLYLSCKVTC--TGCROQEQFRDRSGNVCNCGCPGMELSKGCGFYGEDAQCVT 74
DB 18 LCVLL--VPLQVTLQVTPPTQERHYEHLGRC--CSRCEPGKYLSSKC--TPTSDSVCLP 71
QY 75 CRLHREKEDWGQ-KC---KPC-LDCAVV-----NRFQKANCAT-----SDAIC----- 114
DB 72 CGPDEYLDTWNEEDKCLLHKVCDAGKALVADPGNHTAPRCACCTAGYHWNSDCECCRN 131
QY 115 GDCLPGFYRKTKLVGFQDMECVPC-----GDPPPPY-----EPH----- 148
DB 132 TECAPGFGAHPQLNKNQVCTPCLLGFSDVFSSTDKCPWNTNCTLLGKLEAHQGTES 191
QY 149 ---CASKVNLVIATPASSPRODTALAAVICSALATVLLALLILCVYCKRQFMKKPSWS 205
DB 192 DVMCSWTLRRPPKQAQVLPSLI--VILLFISVVVVAIIFGVYRKGKGTALANLN 249
QY 206 -LRSQDIQYNGSELSCLDRLPOLHEYAHRAAC---QCRDSDVQTCGPVRLPLPSMCCBEACS 261
DB 250 WYNDACSSLSGNKESSGDR-----CAGSHSATSSQVECEGILL---MTREKMW 296
QY 262 PNPATLGCYVHSAAS-----LQARNAGPAGEMVPT-----FFGS 295
DB 297 PEDGAGVCGPVCAAGGPAEVRDSTRFTLVSEVETQGLSRKPTDEYTDPRFSQFSTGS 356
QY 296 LTQSIGGFSDAWPLMNPMD---GGDNISFC-----DSYPELT 330
DB 357 LL--LIQQGSKSLPPQEPLEVEGENDLSQCFTGTGTSTVDSECCDTEPPSPRTDSMP--V 412
QY 331 GEDIHSLNPELSSTSL----DSNSSQDLVG-GAVPVQSH 365
DB 413 SPEKH-LTKEIEGDSCLPWWVSSNSTDGYTGSNGTPEGDH 451

RESULT 15
US-08-995-659-15
; Sequence 15, Application US/08995659
; Patent No. 6242213
; GENERAL INFORMATION:
; APPLICANT: Anderson, Dirk M.
; APPLICANT: Galibert, Laurent
; APPLICANT: Maraskovsky, Eugene
; TITLE OF INVENTION: Ligand for Receptor Activator of NF-kappaB
; NUMBER OF SEQUENCES: 19
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation, Law Department
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: Apple Power Macintosh
; OPERATING SYSTEM: Apple Operating System 7.5.5
; SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/995,659
; FILING DATE: 22 DECEMBER 1997
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: USSN 60/064,671
; FILING DATE: 14 OCTOBER 1997
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: USSN 08/813,509
; FILING DATE: 07 MARCH 1997
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: USSN 08/772,330
; FILING DATE: 23 DECEMBER 1996
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia Anne
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2852-A
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206)587-0430
; TELEFAX: (206)233-0644
; INFORMATION FOR SEQ ID NO: 15:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 625 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-995-659-15

Search completed: April 10, 2002, 17:16:25
Job time: 88 sec

Query Match 6.0%; Score 136; DB 4; Length 625;
Best Local Similarity 22.2%; Pred. No. 0.00023;
Matches 102; Conservative 51; Mismatches 171; Indels 136; Gaps 27;
QY 16 LVLVLLGYSCKVTCE-TGDCRQGEFRDRSGNCVPCNCGFMELSKCEGFGYGEDAQCVT 74
DB 18 LCVLL--VPLQVTLQVTPPTQERHYHLGR--CSRCEPGRKYLSSKC--TPTSDSVCLP 71
QY 75 CRLHREFKEDWGFQ-KC---KPC-LDCAVV-----NRFQKANCST-----SDAIC---- 114
DB 72 CGPDEYLDTWNEEDKCLLHKVCDAGALVAVDPGNHTAPRRCACTAGYHWNDSCECCRN 131
QY 115 GDCPLGFGYRKTLVGFQDMCEVCPC-----GDPpppy-----EPH----- 148
DB 132 TECAPCGGAQHPQLANKDIVCTCLLGFSDVFSSDCKPWTNCTLLGKLEAHQGTES 191
QY 149 ---CASKVNLVKTASTASSPRDTALAAVTCSSALATVLLALLLCVLYCKRQFMKKPSWS 205

Db 192 DVVSSMTLRRPPKEAQAYLPSLI--VLLLFISVVVVAALIFGVYVYRKGGKALTANLWN 249
QY 206 -LRSQDIQYNGSELSCLDRLPOLHEYAHRACC---QCRRDSVQTCGPVRLLPSCCEEACS 261
DB 250 WYNDACSSLSGNKESGDR-----CAGSHSATSSQOEVCGEILL---MTREKMY 296
QY 262 PNPATLGCQGVHSAAS-----LQARNAGPAGEMVPT-----FFGS 295
DB 297 PEDGAGVCGPVCAAGGFWAEVRDSRTFTLYSEVETOGDLRSRKIPTEDYTDPSQBSTGS 356
QY 296 LTQSGICEFSDAWPLMQNPM---GGDNISFC-----DSYPELT 330
DB 357 LL--LIQOGSKSIPPFQEPLEVGENDLSQCFTGTSTVDSECGDFTPEPSTRDTSMP--V 412
QY 331 GEDIHSLNPELESSTSL-----DSNSSQDLVG-CAVPVQSH 365
DB 413 SPEKH-LTRKEIEGDSCLPMWVSSNSTDGYTSGNTGCEDH 451

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